

(19) World Intellectual Property Organization  
International Bureau



(43) International Publication Date  
5 April 2001 (05.04.2001)

PCT

(10) International Publication Number  
WO 01/22972 A2

(51) International Patent Classification<sup>7</sup>: A61K 31/7088, 39/39, 48/00, 35/12, A23L 1/30, A61P 37/04 // (A61K 31/7088, 31:00) (A61K 31/7088, 38:00) (A61K 31/7088, 39:395)

(21) International Application Number: PCT/US00/26383

(22) International Filing Date:  
25 September 2000 (25.09.2000)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:  
60/156,113 25 September 1999 (25.09.1999) US  
60/156,135 27 September 1999 (27.09.1999) US  
60/227,436 23 August 2000 (23.08.2000) US

(71) Applicants: UNIVERSITY OF IOWA RESEARCH FOUNDATION [US/US]; 214 Technology Innovation Center, Oakdale Research Campus, Iowa City, IA 52242 (US). COLEY PHARMACEUTICAL GMBH [DE/DE]; Elisabeth-Selbert-Strasse 9, D-40764 Langenfeld (DE).

(72) Inventors: KRIEG, Arthur, M.; University of Iowa, Dept. of Internal Medicine, 540 EM RB, Iowa City, IA 52242 (US). SCHETTER, Christian; Coley Pharmaceutical Group GmbH, Qiagen GmbH, Max-Volmer Strabe 4, D-40724 Hilden (DE). VOLLMER, Jörg; Coley Pharmaceutical Group GmbH, Qiagen GmbH, Max-Volmer Strabe 4, D-40724 Hilden (DE).

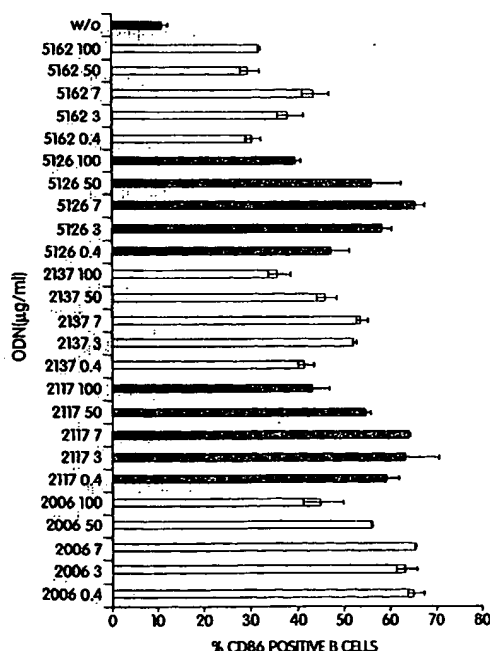
(74) Agent: LOCKHART, Helen, C.; Wolf, Greenfield & Sacks, P.C., 600 Atlantic Avenue, Boston, MA 02210 (US).

(81) Designated States (*national*): AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, UZ, VN, YU, ZA, ZW.

(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

[Continued on next page]

(54) Title: IMMUNOSTIMULATORY NUCLEIC ACIDS



(57) Abstract: The invention relates to immunostimulatory nucleic acid compositions and methods of using the compositions. The T-rich nucleic acids contain poly T sequences and/or have greater than 25 % T nucleotide residues. The TG nucleic acids have TG dinucleotides. The C-rich nucleic acids have at least one poly-C region and/or greater than 50 % c nucleotides. These immunostimulatory nucleic acids function in a similar manner to nucleic acids containing CpG motifs. The invention also encompasses preferred CpG nucleic acids.

WO 01/22972 A2



**Published:**

— Without international search report and to be republished upon receipt of that report.

*For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.*

## **IMMUNOSTIMULATORY NUCLEIC ACIDS**

### **BACKGROUND OF THE INVENTION**

Bacterial DNA has immune stimulatory effects to activate B cells and natural  
5 killer cells, but vertebrate DNA does not (Tokunaga, T., et al., 1988. *Jpn. J. Cancer Res.*  
79:682-686; Tokunaga, T., et al., 1984, *JNCI* 72:955-962; Messina, J.P., et al., 1991, *J.*  
*Immunol.* 147:1759-1764; and reviewed in Krieg, 1998, In: *Applied Oligonucleotide*  
*Technology*, C.A. Stein and A.M. Krieg, (Eds.), John Wiley and Sons, Inc., New York,  
NY, pp. 431-448). It is now understood that these immune stimulatory effects of  
10 bacterial DNA are a result of the presence of unmethylated CpG dinucleotides in  
particular base contexts (CpG motifs), which are common in bacterial DNA, but  
methylated and underrepresented in vertebrate DNA (Krieg et al, 1995 *Nature* 374:546-  
549; Krieg, 1999 *Biochim. Biophys. Acta* 93321:1-10). The immune stimulatory effects  
of bacterial DNA can be mimicked with synthetic oligodeoxynucleotides (ODN)  
15 containing these CpG motifs. Such CpG ODN have highly stimulatory effects on human  
and murine leukocytes, inducing B cell proliferation; cytokine and immunoglobulin  
secretion; natural killer (NK) cell lytic activity and IFN- $\gamma$  secretion; and activation of  
dendritic cells (DCs) and other antigen presenting cells to express costimulatory  
molecules and secrete cytokines, especially the Th1-like cytokines that are important in  
20 promoting the development of Th1-like T cell responses. These immune stimulatory  
effects of native phosphodiester backbone CpG ODN are highly CpG specific in that the  
effects are essentially abolished if the CpG motif is methylated, changed to a GpC, or  
otherwise eliminated or altered (Krieg et al, 1995 *Nature* 374:546-549; Hartmann et al,  
1999 *Proc. Natl. Acad. Sci USA* 96:9305-10). Phosphodiester CpG ODN can be  
25 formulated in lipids, alum, or other types of vehicles with depot properties or improved  
cell uptake in order to enhance the immune stimulatory effects (Yamamoto et al, 1994  
*Microbiol. Immunol.* 38:831-836; Gramzinski et al, 1998 *Mol. Med.* 4:109-118).

In early studies, it was thought that the immune stimulatory CpG motif followed  
the formula purine-purine-CpG-pyrimidine-pyrimidine (Krieg et al, 1995 *Nature*  
30 374:546-549; Pisetsky, 1996 *J. Immunol.* 156:421-423; Hacker et al., 1998 *EMBO J.*  
17:6230-6240; Lipford et al, 1998 *Trends in Microbiol.* 6:496-500). However, it is now  
clear that mouse lymphocytes respond quite well to phosphodiester CpG motifs that do

- 2 -

not follow this "formula" (Yi et al., 1998 J. Immunol. 160:5898-5906) and the same is true of human B cells and dendritic cells (Hartmann et al, 1999 Proc. Natl. Acad. Sci USA 96:9305-10; Liang, 1996 J. Clin. Invest. 98:1119-1129).

Several past investigators have looked at whether the nucleotide content of ODN  
5 may have effects independently of the sequence of the ODN. Interestingly, antisense  
ODN have been found to be generally enriched in the content of GG, CCC, CC, CAC,  
and CG sequences, while having reduced frequency of TT or TCC nucleotide sequences  
compared to what would be expected if base usage were random (Smetsers et al., 1996  
Antisense Nucleic Acid Drug Develop. 6:63-67). This raised the possibility that the  
10 over-represented sequences may comprise preferred targeting elements for antisense  
oligonucleotides or visa versa. One reason to avoid the use of thymidine-rich ODN for  
antisense experiments is that degradation of the ODN by nucleases present in cells  
releases free thymidine which competes with <sup>3</sup>H-thymidine which is frequently used in  
experiments to assess cell proliferation (Matson et al., 1992 Antisense Research and  
15 Development 2:325-330).

#### SUMMARY OF THE INVENTION

The present invention relates in part to pyrimidine rich (Py-rich) and in some  
embodiments thymidine (T) rich immunostimulatory nucleic acids which do not require  
the presence of a CpG motif. The present invention also relates in part to the discovery  
20 that nucleic acids which contain a TG dinucleotide motif are also immunostimulatory.  
The invention is based in part on the unexpected finding that nucleic acid sequences  
which do not contain CpG motifs are immunostimulatory. It was discovered upon  
analysis of the immune stimulation properties of many nucleic acid sequences that these  
sequences may be Py-rich e.g., T-rich or that they may contain TG motifs. It was also  
25 discovered that these sequences preferentially activate non-rodent immune cells. The  
Py-rich and TG sequences are only minimally immunostimulatory with respect to rodent  
immune cells, compared to non-rodent immune cells. Thus, it is possible according to  
the methods of the invention to induce an immune response in a non-rodent subject by  
administering Py-rich or TG immunostimulatory nucleic acids. The Py-rich and TG  
30 immunostimulatory nucleic acids of the invention may optionally include CpG motifs.  
These findings have important implications for the clinical development of  
immunostimulatory CpG containing and non-CpG containing nucleic acids.



- 3 -

In one aspect the invention is a pharmaceutical composition comprising an effective amount for stimulating an immune response of isolated Py-rich or TG immunostimulatory nucleic acids, and a pharmaceutically acceptable carrier. In other aspects the invention is a composition of matter, comprising an isolated Py-rich or TG immunostimulatory nucleic acid. In other embodiments, the immunostimulatory nucleic acid may be T-rich. In still other embodiments, the immunostimulatory nucleic acid may be T-rich and also have at least one TG motif.

Preferably the Py-rich nucleic acid is a T-rich nucleic acid. In some embodiments the T-rich immunostimulatory nucleic acid is a poly T nucleic acid comprising 5' TTTT 3'. In yet other embodiments the poly T nucleic acid comprises 5' X<sub>1</sub> X<sub>2</sub> TTTT X<sub>3</sub> X<sub>4</sub> 3' wherein X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub> and X<sub>4</sub> are nucleotides. In some embodiments X<sub>1</sub>X<sub>2</sub> is TT and/or X<sub>3</sub>X<sub>4</sub> is TT. In other embodiments X<sub>1</sub>X<sub>2</sub> is selected from the group consisting of TA, TG, TC, AT, AA, AG, AC, CT, CC, CA, CG, GT, GG, GA, and GC; and/or X<sub>3</sub>X<sub>4</sub> is selected from the group consisting of TA, TG, TC, AT, AA, AG, AC, CT, CC, CA, CG, GT, GG, GA, and GC.

The T-rich immunostimulatory nucleic acid may have only a single poly T motif or it may have a plurality of poly T nucleic acid motifs. In some embodiments the T-rich immunostimulatory nucleic acid comprises at least 2, at least 3, at least 4, at least 5, at least 6, at least 7, or at least 8 T motifs. In other embodiments it comprises at least 2, at least 3, at least 4, at least 5, at least 6, at least 7, or at least 8 CpG motifs. In preferred embodiments the plurality of CpG motifs and poly T motifs are interspersed.

In yet other embodiments at least one of the plurality of poly T motifs comprises at least 3, at least 4, at least 5, at least 6, at least 7, at least 8, or at least 9 contiguous T nucleotide residues. In other embodiments the plurality of poly T motifs is at least 3 motifs and wherein at least 3 motifs each comprises at least 3 contiguous T nucleotide residues or the plurality of poly T motifs is at least 4 motifs and wherein the at least 4 motifs each comprises at least 3 contiguous T nucleotide residues.

In some cases the T-rich immunostimulatory nucleic acid may be free of poly T motifs but may rather comprise a nucleotide composition of greater than 25% T. In other embodiments the T-rich immunostimulatory nucleic acids have poly T motifs and also comprise a nucleotide composition of greater than 25% T. In preferred embodiments the T-rich immunostimulatory nucleic acid comprises a nucleotide composition of greater

- 4 -

than 35% T, greater than 40% T, greater than 50% T, greater than 60% T, greater than 80% T, or greater than 90% T nucleotide residues. In important embodiments, the nucleic acid is at least 50% T.

The T-rich and TG immunostimulatory nucleic acids can have any length greater than 7 nucleotides, but in some embodiments can be between 8 and 100 nucleotide residues in length. In preferred embodiments the T-rich immunostimulatory nucleic acid comprises at least 20 nucleotides, at least 24 nucleotides, at least 27, nucleotides, or at least 30 nucleotides. In preferred embodiments, the TG immunostimulatory nucleic acid is between 15 and 25 nucleotides in length. The T-rich and TG immunostimulatory nucleic acids may be single stranded or double stranded.

In one preferred embodiment, the immunostimulatory nucleic acid has a T-rich region located in the middle of its length (i.e., an approximately equal number of nucleotides flank the T-rich region on the 5' and 3' ends).

The T rich nucleic acid in some embodiments is selected from the group consisting of SEQ ID NO: 59-63, 73-75, 142, 215, 226, 241, 267-269, 282, 301, 304, 330, 342, 358, 370-372, 393, 433, 471, 479, 486, 491, 497, 503, 556-558, 567, 694, 793-794, 797, 833, 852, 861, 867, 868, 882, 886, 905, 907, 908, and 910-913. In other embodiments the T rich nucleic acids are sequence selected from the group consisting of SEQ ID NO: 64, 98, 112, 146, 185, 204, 208, 214, 224, 233, 244, 246, 247, 258, 262, 263, 265, 270-273, 300, 305, 316, 317, 343, 344, 350, 352, 354, 374, 376, 392, 407, 411-413, 429-432, 434, 435, 443, 474, 475, 498-501, 518, 687, 692, 693, 804, 862, 883, 884, 888, 890, and 891.

In other embodiments the Py-rich immunostimulatory nucleic acid is a C-rich nucleic acid. An immunostimulatory C-rich nucleic acid is a nucleic acid including at least one and preferably at least 2 poly-C regions or which includes 50% or greater C nucleotides.

The Py-rich and TG immunostimulatory nucleic acids may include one or more CpG motifs. The motifs may be methylated or unmethylated. In other embodiments the Py-rich and TG immunostimulatory nucleic acids are free of one or more CpG dinucleotides.

In other embodiments the Py-rich and TG immunostimulatory nucleic acids also include poly-A, poly G, and/or poly C motifs. In yet other embodiments the Py-rich or

- 5 -

TG immunostimulatory nucleic acid is free of two poly C sequences of at least 3 contiguous C nucleotide residues or is free of two poly A sequences of at least 3 contiguous A nucleotide residues. In other embodiments the Py-rich or TG immunostimulatory nucleic acid comprises a nucleotide composition of greater than 25% C or greater than 25% A. In yet other embodiments the Py-rich or TG immunostimulatory nucleic acid is free of poly-C sequences, poly-G sequences or poly-A sequences.

A poly G nucleic acid in some embodiments is selected from the group consisting of SEQ ID NO: 5, 6, 73, 215, 267-269, 276, 282, 288, 297-299, 355, 359, 386, 387, 444, 476, 531, 557-559, 733, 768, 795, 796, 914-925, 928-931, 933-936, and 938. In other embodiments the poly G nucleic acid includes a sequence selected from the group consisting of SEQ ID NO: 67, 80-82, 141, 147, 148, 173, 178, 183, 185, 214, 224, 264, 265, 315, 329, 434, 435, 475, 519, 521-524, 526, 527, 535, 554, 565, 609, 628, 660, 661, 662, 725, 767, 825, 856, 857, 876, 892, 909, 926, 927, 932, and 937.

According to another aspect of the invention, the immunostimulatory nucleic acids may be defined as those which possess a TG motif, herein referred to as TG immunostimulatory nucleic acids. The TG nucleic acid in one embodiment contains at least one TG dinucleotide having a sequence including at least the following formula:  $5'N_1X_1TGX_2N_23'$ . In related embodiments,  $N_1$  is a nucleic acid sequence composed of a number of nucleotides ranging from  $(11-N_2)$  to  $(21-N_2)$  and  $N_2$  is a nucleic acid sequence composed of a number of nucleotides ranging from  $(11-N_1)$  to  $(21-N_1)$ . In a preferred embodiment,  $X_2$  is thymidine.

In other embodiments, the TG nucleic acid has at least the following formula:  $5'X_1X_2TGX_3X_43'$ . In yet another embodiment, the TG nucleic acid comprises the following sequence:  $5'N_1X_1X_2TGX_3X_4N_23'$ . In a related embodiment,  $N_1$  is a nucleic acid sequence composed of a number of nucleotides ranging from  $(9-N_2)$  to  $(19-N_2)$  and  $N_2$  is a nucleic acid sequence composed of a number of nucleotides ranging from  $(9-N_1)$  to  $(19-N_1)$ . In one preferred embodiment,  $X_3$  is thymidine.  $X_1X_2$  are nucleotides which may be selected from the group consisting of GT, GG, GA, AA, AT, AG, CT, CA, CG, TA and TT, and  $X_3X_4$  are nucleotides which may be selected from the group consisting of TT, CT, AT, AG, CG, TC, AC, CC, TA, AA, and CA. In some preferred embodiments,  $X_3$  is a thymidine. In important embodiments,  $X_3X_4$  are nucleotides

- 6 -

selected from the group consisting of TT, TC, TA and TG. In other embodiments  $X_1X_2$  are GA or GT and  $X_3X_4$  are TT. In yet other embodiments  $X_1$  or  $X_2$  or both are purines and  $X_3$  or  $X_4$  or both are pyrimidines or  $X_1X_2$  are GpA and  $X_3$  or  $X_4$  or both are pyrimidines. In one embodiment  $X_2$  is a T and  $X_3$  is a pyrimidine.

5           In one embodiment the 5'  $X_1 X_2 T G X_3 X_4$  3' sequence of the TG nucleic acid or the entire length or some fragment thereof of the TG nucleic acid is a non-palindromic sequence, and in other embodiments it is a palindromic sequence.

In some preferred embodiments, the TG nucleic acid is also T-rich.

10           The Py-rich and TG immunostimulatory nucleic acids in some embodiments have a nucleotide backbone which includes at least one backbone modification, such as a phosphorothioate modification. The nucleotide backbone may be chimeric, or preferably the nucleotide backbone is entirely modified. In one preferred embodiment, the immunostimulatory nucleic acid has a poly T motif and a phosphorothioate backbone.

15           In another aspect the invention is a composition of an immunostimulatory nucleic acid, in the form of a Py-rich or a TG nucleic acid, and an antigen, wherein the nucleic acid is free of unmethylated CpG motifs.

20           Another composition of the invention is a Py-rich or TG immunostimulatory nucleic acid and an anti-microbial agent, wherein the Py-rich or TG nucleic acid is free of unmethylated CpG motifs. Preferably the anti-microbial agent is selected from the group consisting of an anti-viral agent, an anti-parasitic agent, an anti-bacterial agent and an anti-fungal agent.

25           A composition of a sustained release device including a Py-rich and/or TG immunostimulatory nucleic acid, wherein the Py-rich and/or TG nucleic acid is free of unmethylated CpG motifs, is provided according to another aspect of the invention.

30           The invention also includes nutritional supplements of a Py-rich or TG immunostimulatory nucleic acid in a delivery device selected from the group consisting of a capsule, a pill, and a sublingual tablet, wherein the Py-rich or TG nucleic acid is free of unmethylated CpG motifs.

          It should be understood that when it is useful to administer a Py-rich e.g., poly T, T-rich, C-rich, or TG oligonucleotide and a CpG oligonucleotide, it may also be desirable to co-administer a Py-rich or a TG oligonucleotide together with a physically separate CpG, Py-rich or TG oligonucleotide. Alternatively, the CpG, Py-rich or TG

- 7 -

motif may be present on the same contiguous nucleic acid as the Py-rich or TG oligonucleotide. In yet a further embodiment, all or some combination of Py-rich, TG and CpG nucleic acids may be co-administered either on separate nucleic acids or in the same nucleic acid molecule. By co-administer it is intended that the nucleic acids be administered close enough in time to one another to achieve a combined benefit of both oligonucleotides, preferably more than the benefit achieved by administering each of the oligonucleotides alone at the same dose.

CpG oligonucleotides have, in general, the formula 5'X<sub>1</sub>X<sub>2</sub>CGX<sub>3</sub>X<sub>4</sub>3', wherein X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub> and X<sub>4</sub> are nucleotides and wherein at least the C of CpG is unmethylated.

Preferred CpG oligonucleotides are 8-100 nucleotides in length and have modified back bones. Particular structures are detailed in the published PCT applications, U.S. applications and references cited herein, the disclosures of which are incorporated herein in their entirety. In one embodiment, the CpG oligonucleotide is free of poly T and TG motifs and is not T-rich.

In other embodiments, the CpG oligonucleotide has a sequence selected from the group consisting of SEQ ID NO: 1, 3, 4, 14-16, 18-24, 28, 29, 33-46, 49, 50, 52-56, 58, 64-67, 69, 71, 72, 76-87, 90, 91, 93, 94, 96, 98, 102-124, 126-128, 131-133, 136-141, 146-150, 152-153, 155-171, 173-178, 180-186, 188-198, 201, 203-214, 216-220, 223, 224, 227-240, 242-256, 258, 260-265, 270-273, 275, 277-281, 286-287, 292, 295-296, 300, 302, 305-307, 309-312, 314-317, 320-327, 329, 335, 337-341, 343-352, 354, 357, 361-365, 367-369, 373-376, 378-385, 388-392, 394, 395, 399, 401-404, 406-426, 429-433, 434-437, 439, 441-443, 445, 447, 448, 450, 453-456, 460-464, 466-469, 472-475, 477, 478, 480, 483-485, 488, 489, 492, 493, 495-502, 504-505, 507-509, 511, 513-529, 532-541, 543-555, 564-566, 568-576, 578, 580, 599, 601-605, 607-611, 613-615, 617, 619-622, 625-646, 648-650, 653-664, 666-697, 699-706, 708, 709, 711-716, 718-732, 736, 737, 739-744, 746, 747, 749-761, 763, 766-767, 769, 772-779, 781-783, 785-786, 790-792, 798-799, 804-808, 810, 815, 817, 818, 820-832, 835-846, 849-850, 855-859, 862, 865, 872, 874-877, 879-881, 883-885, 888-904, and 909-913.

In another embodiment, the Py-rich or TG oligonucleotide is free of a CpG motifs. This embodiment of the invention also involves pharmaceutical compositions and kits which contain both a CpG oligonucleotide (which can be free of poly T and TG motifs and not be T-rich) and a Py-rich and/or TG oligonucleotide physically separate

- 8 -

from the CpG oligonucleotide. The pharmaceutical preparations are in effective amounts and typically include pharmaceutically acceptable carriers, all as set forth in detail herein with respect to Py-rich and TG oligonucleotides. The kits include at least one container containing an oligonucleotide which is a Py-rich or TG oligonucleotide (or some  
5 combination thereof). The same container, or in other embodiments, a second container, may contain an oligonucleotide with a CpG motif, which may be free of Py-rich and/or TG motifs. The kit also contains instructions for administering the oligonucleotides to a subject. The kits also may include a container containing a solvent or a diluent.

In summary, as if fully recited herein, a CpG oligonucleotide physically separate  
10 from the Py-rich or TG oligonucleotide can be used together with the Py-rich or TG oligonucleotides in the methods, compositions and products described above.

The invention relates in other aspects to immunostimulatory oligonucleotides which have chimeric backbones and which do not require the presence of a CpG motif. The invention is based in part on the discovery that nucleic acid sequences which did not  
15 contain CpG motifs were immunostimulatory, and that those which have chimeric backbones have unexpectedly enhanced immune stimulating properties. Thus the invention in one aspect relates to a composition of an oligonucleotide having a formula:  $5' Y_1 N_1 Z N_2 Y_2 3'$ , wherein  $Y_1$  and  $Y_2$  are, independent of one another, nucleic acid molecules having between 1 and 10 nucleotides, wherein  $Y_1$  includes at least one  
20 modified internucleotide linkage and  $Y_2$  includes at least one modified internucleotide linkage and wherein  $N_1$  and  $N_2$  are nucleic acid molecules, each independent of one another, having between 0 and 5 nucleotides, but wherein  $N_1 Z N_2$  has at least 6 nucleotides in total and wherein the nucleotides of  $N_1 Z N_2$  have a phosphodiester backbone, and wherein  $Z$  is an immunostimulatory nucleic acid motif but does not  
25 include a CG. In one embodiment  $Z$  is a nucleic acid sequence selected from the group consisting of TTTT, TG, and a sequence wherein at least 50% of the bases of the sequence are Ts.

In some embodiments  $Y_1$  and/or  $Y_2$  have between 3 and 8 nucleotides. In other embodiments  $Y_1$  and/or  $Y_2$  are comprised of at least three Gs, at least four Gs, least  
30 seven Gs, or all Gs. In other embodiments  $Y_1$  and/or  $Y_2$  are selected from the group consisting of TCGTCG, TCGTCGT, and TCGTCGTT (SEQ ID NO:1145). In yet other

- 9 -

embodiments  $Y_1$  and/or  $Y_2$  include at least one, two, three, four, or five poly-A, poly-T, or poly-C sequences.

The center nucleotides ( $N_1ZN_2$ ) of the formula  $Y_1N_1ZN_2Y_2$  have phosphodiester internucleotide linkages and  $Y_1$  and  $Y_2$  have at least one modified internucleotide linkage. In some embodiments  $Y_1$  and/or  $Y_2$  have at least two modified internucleotide linkages. In other embodiments  $Y_1$  and/or  $Y_2$  have between two and five modified internucleotide linkages. In yet other embodiments  $Y_1$  has two modified internucleotide linkages and  $Y_2$  has five modified internucleotide linkages or  $Y_1$  has five modified internucleotide linkages and  $Y_2$  has two modified internucleotide linkages. The modified internucleotide linkage, in some embodiments is a phosphorothioate modified linkage, a phosphorodithioate modified linkage or a p-ethoxy modified linkage.

Portions of the formula  $Y_1N_1ZN_2Y_2$  may optionally form a palindrome. Thus, in some embodiments the nucleotides of  $N_1ZN_2$  form a palindrome. In some embodiments the palindrome is not a direct repeat. In yet other embodiments the nucleotides of  $N_1ZN_2$  do not form a palindrome.

According to other embodiments  $N_1ZN_2$  has a sequence of nucleotides selected from the group consisting of GATTTTATCGTC (SEQ ID NO:1098), TCGATTTTTCGA (SEQ ID NO: 1099); TCATTTTATGA (SEQ ID NO: 1100); GTTTTTTACGAC (SEQ ID NO: 1101); TCAATTTTGA (SEQ ID NO: 1102); ACGTTTTTACGT (SEQ ID NO: 1103); TCGTTTTTACGA (SEQ ID NO: 1104); TCGATTTTACGTCGA (SEQ ID NO: 1105); AATTTTAAACGTT (SEQ ID NO: 1106); TCGTTTTTAAACGA (SEQ ID NO: 1107); ACGTTTTTAAACGT (SEQ ID NO: 1108); GATTTTATCGTC (SEQ ID NO: 1109); GACGATTTTTCGTC (SEQ ID NO: 1110); GATTTTAGCTCGTC (SEQ ID NO: 1111); GATTTTACGTC (SEQ ID NO: 1112); ATTTTATCGT (SEQ ID NO: 1113); AACGATTTTTCGTT (SEQ ID NO: 1114); TCACTTTTGTGA (SEQ ID NO: 1115); TCGTATTTA (SEQ ID NO: 1116); ACTTTTGTACCGT (SEQ ID NO: 1117); TCGATTTTTCGACGTCGA (SEQ ID NO: 1118); ACGATTTTTCGT (SEQ ID NO: 1119); GATGATCGTC (SEQ ID NO: 1120); TCGATGTCGA (SEQ ID NO: 1121); TCATGTATGA (SEQ ID NO: 1122); GTGTTACGAC (SEQ ID NO: 1123); TCAATGTTGA (SEQ ID NO: 1124); ACGTGTACGT (SEQ ID NO: 1125); TCGTGTACGA (SEQ ID NO: 1126); TCGATGTACGTCGA (SEQ ID NO: 1127); AATGTAAACGTT (SEQ ID NO: 1128);

- 10 -

TCGTGTAAACGA (SEQ ID NO: 1129); ACGTGTAAACGT (SEQ ID NO: 1130);  
GATGTATCGTC (SEQ ID NO: 1131); GACGATGTCGTC (SEQ ID NO: 1132);  
GATGAGCTCGTC (SEQ ID NO: 1133); GATGTACGTC (SEQ ID NO: 1134);  
ATGATCGT (SEQ ID NO: 1135); AACGATGTCGTT (SEQ ID NO: 1136);  
5 TCACTGGTGA (SEQ ID NO: 1137); TCGTATGA (SEQ ID NO: 1138);  
ACTGGTACCGGT (SEQ ID NO: 1139); TCGATGTCGACGTCGA (SEQ ID NO:  
1140); and ACGATGTCGT (SEQ ID NO: 1141).

The composition may optionally include a pharmaceutical carrier and/or be  
formulated in a delivery device. In some embodiments the delivery device is selected  
10 from the group consisting of cationic lipids, cell permeating proteins, and sustained  
release devices. In one preferred embodiment the sustained release device is a  
biodegradable polymer. In another embodiment the sustained release device is a  
microparticle.

In another aspect the invention is a composition of an immunostimulatory  
15 oligonucleotide having the formula  $Y_1N_1ZN_2Y_2$ , and an antigen.

Another composition of the invention is an immunostimulatory oligonucleotide  
having the formula  $Y_1N_1ZN_2Y_2$ , and an anti-microbial therapeutic agent. Preferably the  
anti-microbial therapeutic agent is selected from the group consisting of an anti-viral  
agent, an anti-parasitic agent, an anti-bacterial agent, or an anti-fungal agent.

20 A composition of a sustained release device including an immunostimulatory  
oligonucleotide having the formula  $Y_1N_1ZN_2Y_2$ , is provided according to another aspect  
of the invention.

The invention also includes nutritional supplements of an immunostimulatory  
oligonucleotide having the formula  $Y_1N_1ZN_2Y_2$ , in a delivery device selected from the  
25 group consisting of a capsule, a sublingual tablet, and a pill.

In another aspect the compositions described above also include an  
immunostimulatory nucleic acid having an unmethylated CG dinucleotide, a TG  
dinucleotide or a Py-rich sequence wherein the immunostimulatory nucleic acid having  
an unmethylated CG dinucleotide, a TG dinucleotide or a Py-rich sequence has a  
30 different sequence than the oligonucleotide comprising 5'  $Y_1N_1ZN_2Y_2$  3'.

In some embodiments the immunostimulatory nucleic acid having an  
unmethylated CG dinucleotide, a TG dinucleotide or a Py-rich sequence has a



completely phosphodiester backbone and in other embodiments the immunostimulatory nucleic acid having an unmethylated CG dinucleotide, a TG dinucleotide or a Py-rich sequence has a modified backbone, which optionally may have internucleotide linkages selected from the group consisting of phosphorothioate, phosphorodithioate, and p-ethoxy.

In one embodiment immunostimulatory nucleic acid having an unmethylated CG dinucleotide has a formula comprising: 5' X<sub>1</sub>X<sub>2</sub>CGX<sub>3</sub>X<sub>4</sub> 3' wherein X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub> and X<sub>4</sub> are nucleotides. In other embodiments the immunostimulatory nucleic acid sequence includes at least the following formula: 5' TCNTX<sub>1</sub>X<sub>2</sub>CGX<sub>3</sub>X<sub>4</sub> 3' wherein N is a nucleic acid sequence composed of from about 0-25 nucleotides, wherein at least one nucleotide has a modified internucleotide linkage, and wherein the nucleic acid has less than or equal to 100 nucleotides. According to some embodiments X<sub>1</sub>X<sub>2</sub> are nucleotides selected from the group consisting of: GT, GG, GA and AA and X<sub>3</sub>X<sub>4</sub> are nucleotides selected from the group consisting of: TT, CT or GT. In a preferred embodiment X<sub>1</sub>X<sub>2</sub> are GA and X<sub>3</sub>X<sub>4</sub> are TT.

In another embodiment the immunostimulatory nucleic acid sequence having an unmethylated CG dinucleotide includes at least one of the following sequences:  
ATCGACTCTCGAGCGTTCTC (SEQ ID No.15); TCCATGTCGGTCCTGCTGAT (SEQ ID No. 32); TCCATGTCGGTZCTGATGCT (SEQ ID No. 31);  
ATCGACTCTCGAGCGTTZTC (SEQ ID No. 18); TCCATGTCGGTCCTGATGCT (SEQ ID No. 28); GGGGGG (SEQ ID No. 12); TCCATGACGGTCCTGATGCT (SEQ ID No. 35); TCCATGGCGGTCCTGATGCT (SEQ ID No. 34);  
TCCATGACGTTCTGATGCT (SEQ ID No. 7); TCCATGTCGTTCTGATGCT (SEQ ID No. 38); GGGGTCAGTCTTGACGGGG (SEQ ID No. 41);  
TCCATGTCGCTCCTGATGCT (SEQ ID No. 37); TCCATGTCGATCCTGATGCT (SEQ ID No. 36); TCCATGCCGGTCCTGATGCT (SEQ ID No. 33);  
TCCATAACGTTCTGATGCT (SEQ ID No. 3); TCCATGACGTTCTGATGCT (SEQ ID No. 7); TCCATGACGTCCCTGATGCT (SEQ ID No 39);  
TCCATCACGTGCCTGATGCT (SEQ ID No. 48); TCCATGACGTTCTGACGTT (SEQ ID No.10); ATGACGTTCTGACGTT (SEQ ID No. 70);  
TCTCCCAGCGCGGCCAT (SEQ ID No. 72); TCCATGTCGTTCTGTCGTT (SEQ ID No. 73); TCCATAGCGTTCTAGCGTT (SEQ ID No. 74);

- 12 -

TCCTGACGTTTCCTGACGTT (SEQ ID No. 76); TCCTGTCGTTTCCTGTCGTT (SEQ ID No. 77); TCCTGTCGTTTCCTGTCGTT (SEQ ID No. 52); TCCTTGTGTCGTTTCCTGTCGTT (SEQ ID No. 121); TCCTGTCGTTTTTTGTCGTT (SEQ ID No. 208); TCGTCGCTGTTGTCGTTTCTT (SEQ ID No. 120);

5 TCCATGCGTTGCGTTGCGTT (SEQ ID No. 81); TCCACGACGTTTTTCGACGTT (SEQ ID No. 82); TCGTCGTTGTCGTTGTCGTT (SEQ ID No. 47); TCGTCGTTTTGTCGTTTTGTCGTT (SEQ ID No. 46); TCGTCGTTGTCGTTTTGTCGTT (SEQ ID No. 49); GCGTGCGTTGTCGTTGTCGTT (SEQ ID No. 56); TGTCGTTTGTGCTTTGTCGTT (SEQ ID No. 48); TGTCGTTGTCGTTGTCGTTGTCGTT (SEQ ID No. 84); TGTCGTTGTCGTTGTCGTT (SEQ ID No. 50); TCGTCGTCGTCGTT (SEQ ID No. 51); and TGTCGTTGTCGTT (SEQ ID No. 85). In another embodiment the immunostimulatory nucleic acid having a Py-rich or TG sequence is a nucleic acid as described above.

15 In another aspect the invention relates to pharmaceutical compositions and kits which contain both an oligonucleotide having the formula  $Y_1N_1ZN_2Y_2$  and a CpG oligonucleotide (which optionally may be free of poly T and TG motifs and not be Py-rich), a Py-rich and/or TG oligonucleotide physically separate from the oligonucleotide having the formula  $Y_1N_1ZN_2Y_2$ . The pharmaceutical preparations are in effective

20 amounts and typically include pharmaceutically acceptable carriers, all as set forth in detail herein. The kits include at least one container containing an oligonucleotide having the formula  $Y_1N_1ZN_2Y_2$ . The same container, or in other embodiments, a second container, may contain an oligonucleotide with a CpG motif, which optionally may be free of Py-rich and/or TG motifs and/or a Py-rich or TG oligonucleotide (or some

25 combination thereof). The kit also contains instructions for administering the oligonucleotides to a subject. The kits also may include a container containing a solvent or a diluent.

In summary, as if fully recited herein, an oligonucleotide having the formula  $Y_1N_1ZN_2Y_2$  which is physically separate from the CpG, Py-rich or TG oligonucleotide

30 can be used together with the CpG, Py-rich, TG oligonucleotides, in the methods, compositions and products described herein.

- 13 -

In another aspect the invention relates to a pharmaceutical composition including at least two oligonucleotides of the invention, wherein the at least two oligonucleotides have different sequences from one another and a pharmaceutically acceptable carrier.

A vaccine formulation is provided according to another aspect of the invention.

- 5 The vaccine includes any of the compositions of the invention in combination with an antigen.

According to another aspect of the invention a method of stimulating an immune response is provided. The method involves administering a Py-rich or a TG immunostimulatory nucleic acid to a non-rodent subject in an amount effective to induce  
10 an immune response in the non-rodent subject. Preferably the Py-rich or TG immunostimulatory nucleic acid is administered orally, locally, in a sustained release device, mucosally to a mucosal surface, systemically, parenterally, or intramuscularly. When the Py-rich or TG immunostimulatory nucleic acid is administered to the mucosal surface it may be delivered in an amount effective for inducing a mucosal immune  
15 response or a systemic immune response. In preferred embodiments the mucosal surface is selected from the group consisting of an oral, nasal, rectal, vaginal, and ocular surface.

In some embodiments the method includes exposing the subject to an antigen wherein the immune response is an antigen-specific immune response. The antigen may be encoded by a nucleic acid vector which can be delivered to the subject. In some  
20 embodiments the antigen is selected from the group consisting of a tumor antigen, a viral antigen, a bacterial antigen, a parasitic antigen and a peptide antigen.

Py-rich and TG immunostimulatory nucleic acids are capable of provoking a broad spectrum of immune response. For instance these immunostimulatory nucleic acids can be used to redirect a Th2 to a Th1 immune response. Py-rich and TG nucleic  
25 acids may also be used to activate an immune cell, such as a leukocyte, a dendritic cell, and an NK cell. The activation can be performed *in vivo*, *in vitro*, or *ex vivo*, i.e., by isolating an immune cell from the subject, contacting the immune cell with an effective amount to activate the immune cell of the Py-rich or TG immunostimulatory nucleic acid and re-administering the activated immune cell to the subject. In some embodiments the  
30 dendritic cell expresses a cancer antigen. The dendritic cell can be exposed to the cancer antigen *ex vivo*.

- 14 -

The immune response produced by Py-rich or TG nucleic acids may also result in induction of cytokine production, e.g., production of IL-6, IL-12, IL-18 TNF, IFN- $\alpha$  and IFN- $\gamma$ .

In still another embodiment, the Py-rich and TG nucleic acids are useful for  
5 treating cancer. The Py-rich and TG nucleic acids are also useful according to other aspects of the invention in preventing cancer (e.g., reducing a risk of developing cancer) in a subject at risk of developing a cancer. The cancer may be selected from the group consisting of biliary tract cancer, breast cancer, cervical cancer, choriocarcinoma, colon cancer, endometrial cancer, gastric cancer, intraepithelial neoplasms, lymphomas, liver  
10 cancer, lung cancer (e.g. small cell and non-small cell), melanoma, neuroblastomas, oral cancer, ovarian cancer, pancreas cancer, prostate cancer, rectal cancer, sarcomas, thyroid cancer, and renal cancer, as well as other carcinomas and sarcomas. In some important embodiments, the cancer is selected from the group consisting of bone cancer, brain and CNS cancer, connective tissue cancer, esophageal cancer, eye cancer, Hodgkin's  
15 lymphoma, larynx cancer, oral cavity cancer, skin cancer, and testicular cancer.

Py-rich and TG nucleic acids may also be used for increasing the responsiveness of a cancer cell to a cancer therapy (e.g., an anti-cancer therapy), optionally when the Py-rich or TG immunostimulatory nucleic acid is administered in conjunction with an anti-cancer therapy. The anti-cancer therapy may be a chemotherapy, a vaccine (e.g., an in  
20 vitro primed dendritic cell vaccine or a cancer antigen vaccine) or an antibody based therapy. This latter therapy may also involve administering an antibody specific for a cell surface antigen of, for example, a cancer cell, wherein the immune response results in antigen dependent cellular cytotoxicity (ADCC). In one embodiment, the antibody may be selected from the group consisting Ributaxin, Herceptin, Quadramet, Panorex,  
25 IDEC-Y2B8, BEC2, C225, Oncolym, SMART M195, ATRAGEN, Ovarex, Bexxar, LDP-03, ior t6, MDX-210, MDX-11, MDX-22, OV103, 3622W94, anti-VEGF, Zenapax, MDX-220, MDX-447, MELIMMUNE-2, MELIMMUNE-1, CEACIDE, Pretarget, NovoMAb-G2, TNT, Gliomab-H, GNI-250, EMD-72000, LymphoCide, CMA 676, Monopharm-C, 4B5, ior egf.r3, ior c5, BABS, anti-FLK-2, MDX-260, ANA Ab,  
30 SMART 1D10 Ab, SMART ABL 364 Ab and ImmuRAIT-CEA.

Thus, according to some aspects of the invention, a subject having cancer or at risk of having a cancer is administered an immunostimulatory nucleic acid and an anti-

cancer therapy. In some embodiments, the anti-cancer therapy is selected from the group consisting of a chemotherapeutic agent, an immunotherapeutic agent and a cancer vaccine. The chemotherapeutic agent may be selected from the group consisting of methotrexate, vincristine, adriamycin, cisplatin, non-sugar containing

5 chloroethylnitrosoureas, 5-fluorouracil, mitomycin C, bleomycin, doxorubicin, dacarbazine, taxol, fragyline, Meglamine GLA, valrubicin, carmustaine and poliferposan, MMI270, BAY 12-9566, RAS farnesyl transferase inhibitor, farnesyl transferase inhibitor, MMP, MTA/LY231514, LY264618/Lometexol, Glamolec, CI-994, TNP-470, Hycamtin/Topotecan, PKC412, Valspodar/PSC833,

10 Novantrone/Mitroxantrone, Metaret/Suramin, Batimastat, E7070, BCH-4556, CS-682, 9-AC, AG3340, AG3433, Incel/VX-710, VX-853, ZD0101, ISI641, ODN 698, TA 2516/Marmistat, BB2516/Marmistat, CDP 845, D2163, PD183805, DX8951f, Lemonal DP 2202, FK 317, Picibanil/OK-432, AD 32/Valrubicin, Metastron/strontium derivative, Temodal/Temozolomide, Evacet/liposomal doxorubicin, Yewtaxan/Paclitaxel,

15 Taxol/Paclitaxel, Xeload/Capecitabine, Furtulon/Doxifluridine, Cyclopax/oral paclitaxel, Oral Taxoid, SPU-077/Cisplatin, HMR 1275/Flavopiridol, CP-358 (774)/EGFR, CP-609 (754)/RAS oncogene inhibitor, BMS-182751/oral platinum, UFT(Tegafur/Uracil), Ergamisol/Levamisole, Eniluracil/776C85/5FU enhancer, Campto/Levamisole, Camptosar/Irinotecan, Tumodex/Ralitrexed, Leustatin/Cladribine, Paxex/Paclitaxel,

20 Doxil/liposomal doxorubicin, Caelyx/liposomal doxorubicin, Fludara/Fludarabine, Pharmarubicin/Epirubicin, DepoCyt, ZD1839, LU 79553/Bis-Naphtalimide, LU 103793/Dolastain, Caetyx/liposomal doxorubicin, Gemzar/Gemcitabine, ZD 0473/Anormed, YM 116, Iodine seeds, CDK4 and CDK2 inhibitors, PARP inhibitors, D4809/Dexifosamide, Ifes/Mesnex/Ifosamide, Vumon/Teniposide,

25 Paraplatin/Carboplatin, Plantinol/cisplatin, Vepeside/Etoposide, ZD 9331, Taxotere/Docetaxel, prodrug of guanine arabinoside, Taxane Analog, nitrosoureas, alkylating agents such as melphelan and cyclophosphamide, Aminoglutethimide, Asparaginase, Busulfan, Carboplatin, Chlorombucil, Cytarabine HCl, Dactinomycin, Daunorubicin HCl, Estramustine phosphate sodium, Etoposide (VP16-213), Floxuridine,

30 Fluorouracil (5-FU), Flutamide, Hydroxyurea (hydroxycarbamide), Ifosfamide, Interferon Alfa-2a, Alfa-2b, Leuprolide acetate (LHRH-releasing factor analogue), Lomustine (CCNU), Mechlorethamine HCl (nitrogen mustard), Mercaptopurine, Mesna,

- 16 -

Mitotane (o.p'-DDD), Mitoxantrone HCl, Octreotide, Plicamycin, Procarbazine HCl, Streptozocin, Tamoxifen citrate, Thioguanine, Thiotepa, Vinblastine sulfate, Amsacrine (m-AMSA), Azacitidine, Erthropoietin, Hexamethylmelamine (HMM), Interleukin 2, Mitoguazone (methyl-GAG; methyl glyoxal bis-guanyldrazone; MGBG), Pentostatin  
5 (2'-deoxycoformycin), Semustine (methyl-CCNU), Teniposide (VM-26) and Vindesine sulfate, but it is not so limited.

The immunotherapeutic agent may be selected from the group consisting of Ributaxin, Herceptin, Quadramet, Panorex, IDEC-Y2B8, BEC2, C225, Oncolym, SMART M195, ATRAGEN, Ovarex, Bexxar, LDP-03, ior t6, MDX-210, MDX-11,  
10 MDX-22, OV103, 3622W94, anti-VEGF, Zenapax, MDX-220, MDX-447, MELIMMUNE-2, MELIMMUNE-1, CEACIDE, Pretarget, NovoMAb-G2, TNT, Gliomab-H, GNI-250, EMD-72000, LymphoCide, CMA 676, Monopharm-C, 4B5, ior egf.r3, ior c5, BABS, anti-FLK-2, MDX-260, ANA Ab, SMART 1D10 Ab, SMART ABL 364 Ab and ImmuRAIT-CEA, but it is not so limited.

The cancer vaccine may be selected from the group consisting of EGF, Anti-idiotypic cancer vaccines, Gp75 antigen, GMK melanoma vaccine, MGv ganglioside conjugate vaccine, Her2/neu, Ovarex, M-Vax, O-Vax, L-Vax, STn-KHL theratope, BLP25 (MUC-1), liposomal idiotypic vaccine, Melacine, peptide antigen vaccines, toxin/antigen vaccines, MVA-based vaccine, PACIS, BCG vaccine, TA-HPV, TA-CIN,  
20 DISC-virus and ImmuCyst/TheraCys, but it is not so limited.

In still another embodiment of the methods directed to preventing or treating cancer, the subject may be further administered interferon- $\alpha$ .

The invention in other aspects relates to methods for preventing disease in a subject. The method involves administering to the subject a Py-rich or a TG  
25 immunostimulatory nucleic acid on a regular basis to promote immune system responsiveness to prevent disease in the subject. Examples of diseases or conditions sought to be prevented using the prophylactic methods of the invention include microbial infections (e.g., sexually transmitted diseases) and anaphylactic shock from food allergies.

30 In other aspects, the invention is a method for inducing an innate immune response by administering to the subject a Py-rich or a TG immunostimulatory nucleic acid in an amount effective for activating an innate immune response.

- 17 -

According to another aspect of the invention a method for treating or preventing a viral or retroviral infection is provided. The method involves administering to a subject having or at risk of having a viral or retroviral infection, an effective amount for treating or preventing the viral or retroviral infection of any of the compositions of the invention.

5 In some embodiments the virus is caused by a hepatitis virus, HIV, hepatitis B, hepatitis C, herpes virus, or papillomavirus.

A method for treating or preventing a bacterial infection is provided according to another aspect of the invention. The method involves administering to a subject having or at risk of having a bacterial infection, an effective amount for treating or preventing  
10 the bacterial infection of any of the compositions of the invention. In one embodiment the bacterial infection is due to an intracellular bacteria.

In another aspect the invention is a method for treating or preventing a parasite infection by administering to a subject having or at risk of having a parasite infection, an effective amount for treating or preventing the parasite infection of any of the  
15 compositions of the invention. In one embodiment the parasite infection is due to an intracellular parasite. In another embodiment the parasite infection is due to a non-helminthic parasite.

In some embodiments the subject is a human and in other embodiments the subject is a non-human vertebrate selected from the group consisting of a dog, cat, horse,  
20 cow, pig, goat, fish, monkey, chicken, and sheep.

In yet another aspect, the invention is a method for treating or preventing asthma, by administering to a subject having or at risk of having asthma, an effective amount for treating or preventing the asthma of any of the compositions of the invention. In one embodiment the asthma is allergic asthma.

25 In another aspect the invention relates to a method for treating or preventing allergy. The method involves administering to a subject having or at risk of having allergy, an effective amount for treating or preventing the allergy of any of the compositions of the invention.

A method for treating or preventing an immune deficiency is provided according  
30 to another aspect of the invention. The method involves administering to a subject having or at risk of an immune deficiency, an effective amount for treating or preventing the immune deficiency of any of the compositions of the invention.

- 18 -

In another aspect the invention relates to a method for inducing a TH1 immune response by administering to a subject any of the compositions of the invention in an effective amount to produce a TH1 immune response.

In one embodiment the methods of the invention involve administering an  
5 oligonucleotide of formula 5' Y<sub>1</sub>N<sub>1</sub>ZN<sub>2</sub>Y<sub>2</sub> 3' and an immunostimulatory nucleic acid having an unmethylated CG dinucleotide a TG dinucleotide or a T-rich sequence. In an embodiment the oligonucleotide comprising 5' Y<sub>1</sub>N<sub>1</sub>ZN<sub>2</sub>Y<sub>2</sub> 3' is administered separately from the immunostimulatory nucleic acid. In some embodiments the oligonucleotide comprising 5' Y<sub>1</sub>N<sub>1</sub>ZN<sub>2</sub>Y<sub>2</sub> 3' and the immunostimulatory nucleic acid are administered  
10 on an alternating weekly schedule and in other embodiments the oligonucleotide comprising 5' Y<sub>1</sub>N<sub>1</sub>ZN<sub>2</sub>Y<sub>2</sub> 3' and the immunostimulatory nucleic acid are administered on an alternating biweekly schedule.

The invention provides in another aspect a composition, comprising an immunostimulatory nucleic acid and an anti-cancer therapy, formulated in a  
15 pharmaceutically-acceptable carrier and in an effective amount to treat a cancer or to reduce the risk of developing a cancer. In important embodiments, the immunostimulatory nucleic acid is selected from the group consisting of a T-rich nucleic acid, a TG nucleic acid and a C-rich nucleic acid.

The invention further provides a kit comprising a first container housing an  
20 immunostimulatory nucleic acid and at least one other container (e.g., a second container) housing an anti-cancer therapy, and instructions for use. In one embodiment, the kit further comprises interferon- $\alpha$ , which may be separately housed in yet another container (e.g., a third container). In an important embodiment, the kit comprises a sustained-release vehicle containing an immunostimulatory nucleic acid, and  
25 at least one container housing an anti-cancer therapy, and instructions for timing of administration of the anti-cancer therapy. The immunostimulatory nucleic acid may be selected from the group consisting of a Py-rich nucleic acid, a TG nucleic acid and a CpG nucleic acid, wherein the CpG nucleic acid has a nucleotide sequence comprising SEQ ID NO: 246.

30 The invention further provides a method for preventing or treating asthma or allergy, comprising administering an immunostimulatory nucleic acid and an asthma/allergy medicament in an effective amount to treat or prevent the asthma or



allergy. In important embodiments, the immunostimulatory nucleic acid is selected from the group consisting of a T-rich nucleic acid, a TG nucleic acid and a C-rich nucleic acid.

In one embodiment the immunostimulatory nucleic acid is a T-rich nucleic acid.

In a related embodiment, the T-rich nucleic acid has a nucleotide sequence selected from  
5 the group consisting of SEQ ID NO: 59-63, 73-75, 142, 215, 226, 241, 267-269, 282, 301, 304, 330, 342, 358, 370-372, 393, 433, 471, 479, 486, 491, 497, 503, 556-558, 567, 694, 793-794, 797, 833, 852, 861, 867, 868, 882, 886, 905, 907, 908, and 910-913. In other embodiments the T-rich nucleic acids are sequence selected from the group consisting of SEQ ID NO: 64, 98, 112, 146, 185, 204, 208, 214, 224, 233, 244, 246, 247,  
10 258, 262, 263, 265, 270-273, 300, 305, 316, 317, 343, 344, 350, 352, 354, 374, 376, 392, 407, 411-413, 429-432, 434, 435, 443, 474, 475, 498-501, 518, 687, 692, 693, 804, 862, 883, 884, 888, 890, and 891.

In yet a further related embodiment, the T-rich nucleic acid is not a TG nucleic acid. In yet still another embodiment, the T-rich nucleic acid is not a CpG nucleic acid.

15 In one embodiment, the immunostimulatory nucleic acid is a TG nucleic acid. In a further related embodiment, the TG nucleic acid is not a T-rich nucleic acid. In another related embodiment, the TG nucleic acid is not a CpG nucleic acid.

In one embodiment, the immunostimulatory nucleic acid is a CpG nucleic acid, wherein the CpG nucleic acid has a nucleotide sequence comprising SEQ ID NO: 246.

20 In another embodiment, the asthma/allergy medicament is a medicament selected from the group consisting of PDE-4 inhibitor, Bronchodilator/beta-2 agonist, K<sup>+</sup> channel opener, VLA-4 antagonist, Neurokin antagonist, TXA2 synthesis inhibitor, Xanthanine, Arachidonic acid antagonist, 5 lipoxygenase inhibitor, Thromboxin A2 receptor antagonist, Thromboxane A2 antagonist, Inhibitor of 5-lipoxygenase activation protein, and  
25 Protease inhibitor, but is not so limited. In some important embodiments, the asthma/allergy medicament is a Bronchodilator/beta-2 agonist selected from the group consisting of salmeterol, salbutamol, terbutaline, D2522/formoterol, fenoterol, and orciprenaline.

In another embodiment, the asthma/allergy medicament is a medicament selected  
30 from the group consisting of Anti-histamines and Prostaglandin inducers. In one embodiment, the anti-histamine is selected from the group consisting of loratidine, cetirizine, buclizine, ceterizine analogues, fexofenadine, terfenadine, desloratadine,

- 20 -

norastemizole, epinastine, ebastine, ebastine, astemizole, levocabastine, azelastine, tranilast, terfenadine, mizolastine, betatastine, CS 560, and HSR 609. In another embodiment, the Prostaglandin inducer is S-5751.

In yet another embodiment, the asthma/allergy medicament is selected from the group consisting of Steroids and Immunomodulators. The immunomodulators may be selected from the group consisting of anti-inflammatory agents, leukotriene antagonists, IL4 muteins, Soluble IL-4 receptors, Immunosuppressants, anti-IL-4 antibodies, IL-4 antagonists, anti-IL-5 antibodies, soluble IL-13 receptor-Fc fusion proteins, anti-IL-9 antibodies, CCR3 antagonists, CCR5 antagonists, VLA-4 inhibitors, and Downregulators of IgE, but are not so limited. In one embodiment, the downregulator of IgE is an anti-IgE.

In another embodiment, the Steroid is selected from the group consisting of beclomethasone, fluticasone, tramcinolone, budesonide, and budesonide. In still a further embodiment, the Immunosuppressant is a Tolerizing peptide vaccine.

In one embodiment, the immunostimulatory nucleic acid is administered concurrently with the asthma/allergy medicament. In another embodiment, the subject is an immunocompromised subject

The immunostimulatory nucleic acids to be administered to a subject in the methods disclosed herein relating to the prevention and treatment of asthma/allergy are as described for other method aspects of the invention.

In another aspect, the invention provides a kit comprising a first container housing an immunostimulatory nucleic acid, and at least another container (e.g., a second container) housing an asthma/allergy medicament, and instructions for use. The immunostimulatory nucleic acid useful in the kit is as described herein. In important embodiments, the immunostimulatory nucleic acid is selected from the group consisting of a T-rich nucleic acid, a TG nucleic acid and a C-rich nucleic acid. In another important embodiment, the kit comprises a sustained-release vehicle containing an immunostimulatory nucleic acid, and at least one container housing an asthma/allergy medicament, and instructions for timing of administration of the asthma/allergy medicament. The asthma/allergy medicament may be selected from the group of asthma/allergy medicaments described in the foregoing methods directed towards the prevention or treatment of asthma/allergy.

In yet another aspect, the invention provides a composition, comprising an immunostimulatory nucleic acid and an asthma/allergy medicament, formulated in a pharmaceutically-acceptable carrier and in an effective amount for preventing or treating an immune response associated with exposure to a mediator of asthma or allergy. The immunostimulatory nucleic acid may be selected from the group of immunostimulatory nucleic acids described for the foregoing methods and compositions. In important embodiments, the immunostimulatory nucleic acid is selected from the group consisting of a T-rich nucleic acid, a TG nucleic acid and a C-rich nucleic acid. The asthma/allergy medicament may be selected from the group consisting of asthma medicaments and allergy medicaments as described in the foregoing methods and compositions.

In still a further aspect, the invention provides a composition comprising an immunostimulatory nucleic acid selected from the group consisting of SEQ ID NO: 95-136, SEQ ID NO: 138-152, SEQ ID NO: 154-222, SEQ ID NO: 224-245, SEQ ID NO: 247-261, SEQ ID NO: 263-299, SEQ ID NO: 301, SEQ ID NO: 303-4109, SEQ ID NO: 414-420, SEQ ID NO: 424, SEQ ID NO: 426-947, SEQ ID NO: 959-1022, SEQ ID NO: 1024-1093, and a pharmaceutically acceptable carrier. Preferably the immunostimulatory nucleic acid is present in the composition in an effective amount. In one embodiment, the immunostimulatory nucleic acid is present in an effective amount to induce an immune response. In another embodiment, the immunostimulatory nucleic acid is present in an effective amount to prevent or treat cancer. In yet a further embodiment, the immunostimulatory nucleic acid is present in an effective amount to prevent or treat asthma/allergy. The invention also provides kits comprising any of the foregoing immunostimulatory nucleic acid compositions, and instructions for use.

In another aspect the invention includes a composition of an immunostimulatory nucleic acid consisting essentially of: 5' M<sub>1</sub>TCGTCGTTM<sub>2</sub> 3' wherein at least one of the Cs is unmethylated, wherein M<sub>1</sub> is a nucleic acid having at least one nucleotide, wherein M<sub>2</sub> is a nucleic acid having between 0 and 50 nucleotides, and wherein the immunostimulatory nucleic acid has less than 100 nucleotides.

In yet other aspects the invention relates to a pharmaceutical composition of an immunostimulatory nucleic acid comprising: 5' TCGTCGTT 3' wherein at least one of the Cs is unmethylated, wherein the immunostimulatory nucleic acid has less than 100 nucleotides and a phosphodiester backbone, and a sustained release device. In some

- 22 -

embodiments the sustained release device is a microparticle. In other embodiments the composition includes an antigen.

Each of the limitations of the invention can encompass various embodiments of the invention. It is, therefore, anticipated that each of the limitations of the invention involving any one element or combinations of elements can be included in each aspect of the invention.

#### BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1A is a histogram of the expression of CD86 (Y-axis) by CD19+ cells following exposure of these cells to the oligonucleotides shown on the X-axis at a concentration of 0.15  $\mu\text{g/ml}$ .

Figure 1B is a histogram of the expression of CD86 (Y-axis) by CD19+ cells following exposure of these cells to the oligonucleotides shown on the X-axis at a concentration of 0.30  $\mu\text{g/ml}$ .

Figure 2 is a graph comparing the abilities of ODN 2137, ODN 2177, ODN 2200 and ODN 2202 to stimulate B cell proliferation at concentrations ranging from 0.2  $\mu\text{g/ml}$  to 20  $\mu\text{g/ml}$ .

Figure 3 is a graph comparing the abilities of ODN 2188, ODN 2189, ODN 2190 and ODN 2182 to stimulate B cell proliferation at concentrations ranging from 0.2  $\mu\text{g/ml}$  to 20  $\mu\text{g/ml}$ .

Fig. 4 is a bar graph depicting dose-dependent B cell activation induced by non-CpG ODN. PBMC of a blood donor were incubated with the indicated concentrations of ODNs 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ ID NO.: 886), 5126 (SEQ ID NO.: 1058) and 5162 (SEQ ID NO.: 1094) and stained with mAb for CD19 (B cell marker) and CD86 (B cell activation marker, B7-2). Expression was measured by flow cytometry.

Fig. 5 is a bar graph depicting stimulation of B cells by a diverse set of non-CpG ODNs. PBMC of one representative donor were stimulated by 0.4 $\mu\text{g/ml}$ , 1.0 $\mu\text{g/ml}$  or 10.0 $\mu\text{g/ml}$  of the following ODNs: 2006 (SEQ ID NO.: 246), 2196 (SEQ ID NO.: 913), 2194 (SEQ ID NO.: 911), 5162 (SEQ ID NO.: 1094), 5163 (SEQ ID NO.: 1095), 5168 (SEQ ID NO.: 1096) and 5169 (SEQ ID NO.: 1097) and expression of the activation marker CD86 (B7-2) on CD19-positive B cells was measured by flow cytometry.

Fig 6 is a bar graph depicting B cell activation by non-CpG ODNs 1982 and 2041. PBMC were incubated with the indicated concentrations of ODN 2006 (SEQ ID NO.: 246), 1982 (SEQ ID NO.: 225) and 2041 (SEQ ID NO.: 282) and B cell activation (expression of the activation marker CD86) was measured by flow cytometry.

5 Fig. 7 is a bar graph depicting NK cells are activated by non-CpG ODNs. PBMC were incubated with 6µg/ml of the following ODNs: 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ ID NO.: 886), 2183 (SEQ ID NO.: 433), 2194 (SEQ ID NO.: 911) and 5126 (SEQ ID NO.: 1058) and stained with mAb for CD3 (T cell marker), CD56 (NK cell marker) and CD69 (early activation marker). Expression of  
10 CD69 on CD56-positive NK cells was measured by flow cytometry.

Fig. 8 is a bar graph depicting NK-mediated cytotoxicity is enhanced by non-CpG ODN. NK-mediated lysis of K-562 target cells was measured after over night incubation of PBMC with 6µg/ml of the ODN 2006 (SEQ ID NO.: 246), 2194 (SEQ ID NO.: 911) and 5126 (SEQ ID NO.: 1058).

15 Fig. 9 is a bar graph depicting NKT cells can be activated by non-CpG ODN. PBMC of one representative donor were incubated with 6µg/ml ODN 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ ID NO.: 886), 2183 (SEQ ID NO.: 433), 2194 (SEQ ID NO.: 911) and 5126 (SEQ ID NO.: 1058) for 24h and activation of NKT cells was measured by flow cytometry after staining of cells with mAb for CD3 (T  
20 cell marker), CD56 (NK cell marker) and CD69 (early activation marker).

Fig. 10 is a bar graph depicting stimulation of monocytes by different CpG and non-CpG ODN. PBMC were incubated with 6µg/ml 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ ID NO.: 886), 2178 (SEQ ID NO.: 428), 2183 (SEQ ID NO.: 433), 2194 (SEQ ID NO.: 911), 5126 (SEQ ID NO.: 1058) and 5163 (SEQ ID NO.:  
25 1095) and stained for CD14 (monocyte marker) and CD80 (B7-1, activation marker). Expression was measured by flow cytometry.

Fig. 11 is a bar graph depicting release of TNFα upon culture of human cells with non-CpG ODN. PBMC were cultured for 24h with or without 6µg/ml of the indicated ODNs or 1µg/ml LPS as positive control and TNFα measured by ELISA.

30 Fig. 12 is a bar graph depicting release of IL-6 after culture with non-CpG ODNs shows the same pattern as for TNFα. PBMC were cultured with the indicated ODNs (1.0µg/ml) and IL-6 was measured in the supernatants by ELISA.

- 24 -

DETAILED DESCRIPTION

The invention in one aspect involves the finding that pyrimidine (Py) rich and preferably thymidine (T) rich nucleic acids as well as nucleic acids that contain TG dinucleotide motifs are effective in mediating immune stimulatory effects. It was known  
5 in the prior art that CpG containing nucleic acids are therapeutic and prophylactic compositions that stimulate the immune system to treat cancer, infectious diseases, allergy, asthma and other disorders and to help protect against opportunistic infections following cancer chemotherapies. The strong yet balanced, cellular and humoral immune responses that result from CpG stimulation reflect the body's own natural  
10 defense system against invading pathogens and cancerous cells. CpG sequences, while relatively rare in human DNA are commonly found in the DNA of infectious organisms such as bacteria. The human immune system has apparently evolved to recognize CpG sequences as an early warning sign of infection, and to initiate an immediate and powerful immune response against invading pathogens without causing adverse reactions  
15 frequently seen with other immune stimulatory agents. Thus CpG containing nucleic acids, relying on this innate immune defense mechanism, can utilize a unique and natural pathway for immune therapy. The effects of CpG nucleic acids on immune modulation were discovered by the inventor of the instant patent application and have been described extensively in co-pending patent applications, such as U.S. Patent Application Serial  
20 Nos: 08/386,063 filed on 02/07/95 (and related PCT US95/01570); 08/738,652 filed on 10/30/96; 08/960,774 filed on 10/30/97 (and related PCT/US97/19791, WO 98/18810); 09/191,170 filed on 11/13/98; 09/030,701 filed on 02/25/98 (and related PCT/US98/03678; 09/082,649 filed on 05/20/98 (and related PCT/US98/10408); 09/325,193 filed on 06/03/99 (and related PCT/US98/04703); 09/286,098 filed on  
25 04/02/99 (and related PCT/US99/07335); 09/306,281 filed on 05/06/99 (and related PCT/US99/09863). The entire contents of each of these patents and patent applications is hereby incorporated by reference.

The findings of the instant invention are applicable to all of the above described uses of CpG containing nucleic acids as well as any other known use for CpG nucleic  
30 acids. The invention involves, in one aspect, the discovery that Py-rich and preferably T-rich and TG nucleic acids have similar immune stimulatory properties to CpG oligonucleotides regardless of whether a CpG motif is present. Thus the invention is

- 25 -

useful for any method for stimulating the immune system using Py-rich or TG nucleic acids. It was also discovered surprisingly according to the invention that chimeric oligonucleotides which lack a CpG motif are immune stimulatory and have many of the same prophylactic and therapeutic activities as a CpG oligonucleotide.

5       A Py-rich nucleic acid is a T-rich or C-rich immunostimulatory nucleic acid. In some embodiments T-rich nucleic acids are preferred. A T-rich nucleic acid is a nucleic acid which includes at least one poly T sequence and/or which has a nucleotide composition of greater than 25% T nucleotide residues. A nucleic acid having a poly-T sequence includes at least four Ts in a row, such as 5'TTTT3'. Preferably the T-rich  
10       nucleic acid includes more than one poly T sequence. In preferred embodiments the T-rich nucleic acid may have 2, 3, 4, etc poly T sequences, such as oligonucleotide #2006 (SEQ ID NO:246). One of the most highly immunostimulatory T-rich oligonucleotides discovered according to the invention is a nucleic acid composed entirely of T nucleotide residues, e.g., oligonucleotide #2183 (SEQ ID NO:433). Other T-rich nucleic acids  
15       according to the invention have a nucleotide composition of greater than 25% T nucleotide residues, but do not necessarily include a poly T sequence. In these T-rich nucleic acids the T nucleotide residues may be separated from one another by other types of nucleotide residues, i.e., G, C, and A. In some embodiments the T-rich nucleic acids have a nucleotide composition of greater than 35%, 40%, 50%, 60%, 70%, 80%, 90%,  
20       and 99%, T nucleotide residues and every integer % in between. Preferably the T-rich nucleic acids have at least one poly T sequence and a nucleotide composition of greater than 25% T nucleotide residues.

      It was discovered according to the invention that the T content of an ODN has a dramatic effect on the immune stimulatory effect of the ODN and that T-rich ODN can  
25       activate multiple human immune cell types in the absence of any CpG motifs. An oligonucleotide having a 3' poly-T region and 2 5'CGs e.g., ODN 2181 (SEQ ID NO:431) is highly immune stimulatory. An oligonucleotide of similar length, ODN 2116 (SEQ ID NO:357) which contains two CG dinucleotides at the 5' end and a poly-C region at the 3' end was also immune stimulatory but to a lesser extent than the T-rich  
30       oligonucleotide using standard experimental conditions. Thus, although C and T have almost identical structures, their effects on the immune properties of an ODN are varied. They both are capable of inducing an immune response but to different extents. Thus

- 26 -

both T-rich and C-rich oligonucleotides are useful according to the invention, but T-rich oligonucleotides are preferred. Furthermore, if the T content of the ODN is reduced by incorporating other bases such as G, A, or C, then the immune stimulatory effects are reduced (ODN #2188 (SEQ ID NO:905), 2190 (SEQ ID NO:907), 2191 (SEQ ID NO:908), and 2193 (SEQ ID NO:910)).

A C-rich nucleic acid is a nucleic acid molecule having at least one or preferably at least two poly-C regions or which is composed of at least 50% C nucleotides. A poly-C region is at least four C residues in a row. Thus a poly-C region is encompassed by the formula 5'CCCC 3'. In some embodiments it is preferred that the poly-C region have the formula 5'CCCCC 3'. Other C-rich nucleic acids according to the invention have a nucleotide composition of greater than 50% C nucleotide residues, but do not necessarily include a poly C sequence. In these C-rich nucleic acids the C nucleotide residues may be separated from one another by other types of nucleotide residues, i.e., G, T, and A. In some embodiments the C-rich nucleic acids have a nucleotide composition of greater than 60%, 70%, 80%, 90%, and 99%, C nucleotide residues and every integer % in between. Preferably the C-rich nucleic acids have at least one poly C sequence and a nucleotide composition of greater than 50% C nucleotide residues, and in some embodiments are also T-rich.

As shown in the Examples, several ODN previously believed to be non-immunostimulatory, including two ODNs SEQ ID NO.: 225 and SEQ ID NO.: 282 previously described to be non-stimulatory and mainly used as control ODNs (Takahashi, T et al 2000. *J. Immunol.* 164:4458) were found to be immunostimulatory. Our experiments, demonstrated that these ODNs can stimulate B cells, although at higher concentrations compared to CpG ODNs (Fig. 6). A long Poly T ODN (30mer) induced, at least in some experiments, comparable strong activation of B cells to one of the strongest CpG ODN activators of B cells. These experiments also revealed the surprising finding that even Poly C ODNs can lead to stimulation of B cells.

Immunostimulation by these ODNs, however, was not limited to human B cells. Different experimental assays clearly demonstrated in addition that monocytes, NK cells and even NKT cells can be activated by such non-CpG ODNs (Fig. 7 – 10). In contrast to Poly T and Poly C sequences, immunostimulation by Poly A sequences (at least for monocytes, B and NK cells) was not achieved. Interestingly it was found that the



- 27 -

introduction of a CpG motif into SEQ ID NO.: 225 enhanced the immunostimulatory activity whereas the elongation with a Poly T stretch did not enhance immunostimulation. This suggests that CpG and T-rich ODN may operate through different mechanisms or pathways. It is also possible that insertion of a poly-T motif  
5 into a different position of SEQ ID NO.: 225 may result in a change in immunostimulatory properties.

A "TG nucleic acid" or a "TG immunostimulatory nucleic acid" as used herein is a nucleic acid containing at least one TpG dinucleotide (thymidine-guanine dinucleotide sequence, i.e. "TG DNA" or DNA containing a 5' thymidine followed by 3' guanosine  
10 and linked by a phosphate bond) and activates a component of the immune system.

In one embodiment the invention provides a TG nucleic acid represented by at least the formula:



wherein  $X_1$  and  $X_2$  are nucleotides and N is any nucleotide and  $N_1$  and  $N_2$  are  
15 nucleic acid sequences composed of any number of N provided that the sum total of  $N_1$  and  $N_2$  is in the range of 11 to 21. As an example, if  $N_1$  is 5, then  $N_2$  may be 6 (leading to a total length for the oligonucleotide of 15 nucleotides). The TG may be located anywhere within the oligonucleotide stretch, including the 5' end, the center and the 3' end. Thus,  $N_1$  may be zero through to 21, inclusive, provided that  $N_2$  is appropriately  
20 chosen to give a sum of  $N_2$  and  $N_1$  equal to 11 through to 21, inclusive. Similarly,  $N_2$  may be zero through to 21, inclusive, provided that the sum total of  $N_1$  and  $N_2$  equals 11 to 21, inclusive. In some embodiments  $X_1$  is adenine, guanine, or thymidine and  $X_2$  is cytosine, adenine, or thymidine. In one preferred embodiment,  $X_2$  is thymidine. In other  
25 herein, the nucleic acid may encompass other motifs, provided it is long enough to do so.

In other embodiments the TG nucleic acid is represented by at least the formula:



wherein  $X_1$ ,  $X_2$ ,  $X_3$ , and  $X_4$  are nucleotides. In some embodiments,  $X_1X_2$  are nucleotides selected from the group consisting of: GpT, GpG, GpA, ApA, ApT, ApG, CpT, CpA, TpA and TpT; and  $X_3X_4$  are nucleotides selected from the group consisting  
30 of: TpT, CpT, ApT, ApG, TpC, ApC, CpC, TpA, ApA, and CpA; N is any nucleotide and  $N_1$  and  $N_2$  are nucleic acid sequences composed of any number of nucleotides

provide that the sum total of  $N_1$  and  $N_2$  is in the range of 9 to 19. In some embodiments,  $X_1X_2$  are GpA or GpT and  $X_3X_4$  are TpT. In other embodiments  $X_1$  or  $X_2$  or both are purines and  $X_3$  or  $X_4$  or both are pyrimidines or  $X_1X_2$  are GpA and  $X_3$  or  $X_4$  or both are pyrimidines. In one preferred embodiment,  $X_3X_4$  are nucleotides selected from the group consisting of: TpT, TpC and TpA.

The immunostimulatory nucleic acid may be any size (i.e., length) provided it is at least 4 nucleotides. In important embodiments, the immunostimulatory nucleic acids have a length in the range of between 6 and 100. In still other embodiments, the length is in the range of between 8 and 35 nucleotides. Preferably, the TG oligonucleotides range in size from 15 to 25 nucleotides.

The size (i.e., the number of nucleotide residues along the length of the nucleic acid) of the immunostimulatory nucleic acid may also contribute to the stimulatory activity of the nucleic acid. It has been discovered, surprisingly that even for highly immune stimulating immunostimulatory nucleic acids, the length of the nucleic acid influences the extent of immunostimulation that can be achieved. It has been demonstrated that increasing the length of a T-rich nucleic acid up to 24 nucleotides causes increased immune stimulation. The experiments presented in the examples demonstrate that when the length of the T-rich nucleic acid is increased from 18 to 27 nucleotides the ability of the nucleic acid to stimulate an immune response is increased significantly (compare ODN #2194, 2183, 2195, and 2196 decreasing in size from 27-18 nucleotides). Increasing the length of the nucleic acid up to 30 nucleotides had a dramatic impact on the biological properties of the nucleic acid but increasing the length beyond 30 nucleotides did not appear to further influence the immune stimulatory effect (e.g., compare ODN 2179 to 2006).

It has been shown that TG nucleic acids ranging in length from 15 to 25 nucleotides in length may exhibit an increased immune stimulation. Thus, in one aspect, the invention provides an oligonucleotide that is 15-27 nucleotides in length (i.e., an oligonucleotide that is 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26 or 27 nucleotides in length) that may be a T-rich nucleic acid or may be a TG nucleic acid, or may be both a T-rich and a TG nucleic acid. In one embodiment, the oligonucleotide is not a T-rich nucleic acid nor is it a TG nucleic acid. In other embodiments, the oligonucleotide does not have a CG motif. The invention similarly provides oligonucleotides that are 15-27

- 29 -

nucleotides in length, oligonucleotides that are 18-25 nucleotides in length, oligonucleotides that are 20-23 nucleotides in length, and oligonucleotides that are 23-25 nucleotides in length. Any of the foregoing embodiments relating to oligonucleotides 15-27 in length also relate to the oligonucleotides of these differing lengths. The invention further embraces the use of any of these foregoing oligonucleotides in the methods recited herein.

Although a maximal level of immune stimulation is achieved with some T-rich nucleic acids when the nucleic acid is 24-30 nucleotide residues in length, as well as with some TG nucleic acids that range from 15 to 25 nucleotides in length, shorter or longer immunostimulatory nucleic acids can also be used according to the methods of the invention. For facilitating uptake into cells immunostimulatory nucleic acids preferably have a minimum length of 6 nucleotide residues. Nucleic acids of any size greater than 6 nucleotides (even many kb long) are capable of inducing an immune response according to the invention if sufficient immunostimulatory motifs are present, since larger nucleic acids are degraded inside of cells. Preferably the immunostimulatory nucleic acids are in the range of between 8 and 100 and in some embodiments T-rich containing immunostimulatory nucleic acids are between 24 and 40 nucleotides in length and TG containing immunostimulatory nucleic acids are between 15 and 25 nucleotides in length.

In one embodiment the T-rich nucleic acid is represented by at least the formula:



wherein  $X_1$ ,  $X_2$ ,  $X_3$ , and  $X_4$  are nucleotides. In one embodiment  $X_1X_2$  is TT and/or  $X_3X_4$  is TT. In another embodiment  $X_1X_2$  are any one of the following nucleotides TA, TG, TC, AT, AA, AG, AC, CT, CC, CA, GT, GG, GA, and GC; and  $X_3X_4$  are any one of the following nucleotides TA, TG, TC, AT, AA, AG, AC, CT, CC, CA, GT, GG, GA, and GC.

In some embodiments it is preferred that the immunostimulatory nucleic acids do not contain poly-C (CCCC), or poly-A (AAAA). In other embodiments it is preferred that the immunostimulatory nucleic acid include poly-C, poly-A, poly-G (GGGG) or multiple GGs. In particular poly-G or multiple GG motifs have dramatic effects on some immunostimulatory nucleic acids. The effect of these non-T sequences depends in part on the status of the nucleic acid backbone. For instance, if the nucleic acid has a

- 30 -

phosphodiester backbone or a chimeric backbone the inclusion of these sequences in the nucleic acid will only have minimal if any effect on the biological activity of the nucleic acid. If the backbone is completely phosphorothioate (or other phosphate modification) or significantly phosphorothioate then the inclusion of these sequences may have more  
5 influence on the biological activity or the kinetics of the biological activity, causing a decrease in potency of the T-rich and TG immunostimulatory nucleic acids.

Although C-rich nucleic acids have been demonstrated to have immune stimulating properties, insertion of Poly-C sequences into a T-rich nucleic acid in a manner that would reduce the relative proportion of T nucleotides in the nucleic acid can  
10 have a negative impact on the nucleic acid. Although applicants are not bound by a proposed mechanism, it is believed that the immune system has developed a mechanism for distinguishing nucleic acids having different nucleotide properties, possibly resulting from different sets of binding proteins which recognize different sequences or specific binding proteins which recognize all the immunostimulatory sequences but with different  
15 affinities. In general nucleic acids including unmethylated CpG motifs are the most immunostimulatory, followed by T-rich nucleic acids, TG nucleic acids and C-rich nucleic acids. This generalization, however, has many exceptions. For instance a strong T-rich nucleic acid like SEQ ID NO.: 886 is more immune stimulatory in some assays than some CpG containing nucleic acids (e.g., a phosphorothioate CpG nucleic acid  
20 containing a single CpG motif).

It has also been discovered that the addition of a poly-A tail to an immunostimulatory nucleic acid can enhance the activity of the nucleic acid. It was discovered that when a highly immune stimulatory CpG nucleic acid (SEQ ID NO.: 246) was modified with the addition of a poly-A tail (AAAAAA) or a poly-T tail (TTTTTT),  
25 the resultant oligonucleotides increased in immune stimulatory activity. The ability of the poly-A tail and the poly-T tail to increase the immunostimulating properties of the oligonucleotide was very similar. SEQ ID NO.: 246 is a T-rich oligonucleotide. It is likely that if poly-A and poly-T tails are added to a nucleic acid which is not T-rich, it would have a bigger impact on the immune stimulating capability of the nucleic acid.  
30 Since the poly-T tail was added to a nucleic acid that was already highly T-rich the immune stimulating properties of the poly-T addition was diluted somewhat, although not completely. This finding has important implications for the use of poly-A regions.

- 31 -

Thus in some embodiments the immunostimulatory nucleic acids include a poly-A region and in other embodiments they do not.

Some of the immunostimulatory nucleic acids of the invention include one or more CG motifs. The presence of CG motifs in the immunostimulatory nucleic acids  
5 also has an influence on the biological activity of the nucleic acids. If the total length of an immunostimulatory nucleic acid is 20 nucleotide residues or less, then CpG motifs are important in determining the immune effect of the nucleic acid, and methylation of these motifs reduces the potency of the immune stimulatory effects of the nucleic acid. If the  
10 length of the immunostimulatory nucleic acid is increased to 24, then the immune stimulatory effects of the nucleic acid become less dependent on the CpG motifs, and are no longer abolished by methylation of the CpG motifs or by their inversion to GC dinucleotides, provided the other immune-stimulatory properties described herein are present.

For example, ODN 2006 (SEQ ID NO:246) is a highly immune stimulatory T-rich  
15 nucleic acid of 24 nucleotide residues in length with four CpG dinucleotides. However, ODN 2117 (SEQ ID NO:358), in which the CpG motifs are methylated is also highly immune stimulatory. ODN 2137 (SEQ ID NO:886), in which the CpG motifs of ODN 2006 are inverted to GpC, and which as a result possesses six TG dinucleotides is also immune stimulatory. The immune stimulatory effects of nucleic acids such as ODN  
20 2117 and 2137 are regulated by their T and TG content. Each of these three nucleic acids is T-rich and ODN 2137 is additionally TG rich. If their T content is reduced by inserting other bases such as A (ODN 2117 (SEQ ID NO:358)) or if their TG content is reduced by substituting TG with AG, then the immune stimulatory effects are somewhat reduced. In another example, a nucleic acid 24 nucleotides in length in which all of the  
25 positions are randomized has only a modest immune stimulatory effect (ODN 2182 (SEQ ID NO:432)). Likewise, a nucleic acid 24 nucleotides in length with other nucleotide compositions have variable immune stimulatory effects, depending on their T content (ODN 2188 (SEQ ID NO:905), 2189 (SEQ ID NO:906), 2190 (SEQ ID NO:907), 2191 (SEQ ID NO:908), 2193 (SEQ ID NO:910), 2183 (SEQ ID NO:433), and  
30 2178 (SEQ ID NO:428)). ODN 2190 which contains TGT motifs is more immune stimulatory than ODN 2202 which possesses TGG motifs. Thus, in some embodiments, TGT motifs are preferred. In still other embodiments, the number of TG motifs is

important in that an increase in the number of TG motifs leads to an increase in immune stimulation. Some preferred TG nucleic acids contain at least three TG motifs.

Examples of CpG nucleic acids include but are not limited to those listed in Table A, such as SEQ ID NO: 1, 3, 4, 14-16, 18-24, 28, 29, 33-46, 49, 50, 52-56, 58, 64-67, 69,  
5 71, 72, 76-87, 90, 91, 93, 94, 96, 98, 102-124, 126-128, 131-133, 136-141, 146-150,  
152-153, 155-171, 173-178, 180-186, 188-198, 201, 203-214, 216-220, 223, 224, 227-  
240, 242-256, 258, 260-265, 270-273, 275, 277-281, 286-287, 292, 295-296, 300, 302,  
305-307, 309-312, 314-317, 320-327, 329, 335, 337-341, 343-352, 354, 357, 361-365,  
367-369, 373-376, 378-385, 388-392, 394, 395, 399, 401-404, 406-426, 429-433, 434-  
10 437, 439, 441-443, 445, 447, 448, 450, 453-456, 460-464, 466-469, 472-475, 477, 478,  
480, 483-485, 488, 489, 492, 493, 495-502, 504-505, 507-509, 511, 513-529, 532-541,  
543-555, 564-566, 568-576, 578, 580, 599, 601-605, 607-611, 613-615, 617, 619-622,  
625-646, 648-650, 653-664, 666-697, 699-706, 708, 709, 711-716, 718-732, 736, 737,  
739-744, 746, 747, 749-761, 763, 766-767, 769, 772-779, 781-783, 785-786, 7900792,  
15 798-799, 804-808, 810, 815, 817, 818, 820-832, 835-846, 849-850, 855-859, 862, 865,  
872, 874-877, 879-881, 883-885, 888-904, and 909-913.

In some embodiments of the invention the immunostimulatory nucleic acids include CpG dinucleotides and in other embodiments the immunostimulatory nucleic acids are free of CpG dinucleotides. The CpG dinucleotides may be methylated or  
20 unmethylated. A nucleic acid containing at least one unmethylated CpG dinucleotide is a nucleic acid molecule which contains an unmethylated cytosine-guanine dinucleotide sequence (i.e. "CpG DNA" or DNA containing an unmethylated 5' cytosine followed by 3' guanosine and linked by a phosphate bond) and activates the immune system. A nucleic acid containing at least one methylated CpG dinucleotide is a nucleic acid which  
25 contains a methylated cytosine-guanine dinucleotide sequence (i.e., a methylated 5' cytosine followed by a 3' guanosine and linked by a phosphate bond).

Examples of T rich nucleic acids that are free of CpG nucleic acids include but are not limited to those listed in Table A, such as SEQ ID NO: 59-63, 73-75, 142, 215, 226, 241, 267-269, 282, 301, 304, 330, 342, 358, 370-372, 393, 433, 471, 479, 486, 491,  
30 497, 503, 556-558, 567, 694, 793-794, 797, 833, 852, 861, 867, 868, 882, 886, 905, 907, 908, and 910-913. Examples of T rich nucleic acids that include CpG nucleic acids include but are not limited to those listed in Table A, such as SEQ ID NO: 64, 98, 112,

- 33 -

146, 185, 204, 208, 214, 224, 233, 244, 246, 247, 258, 262, 263, 265, 270-273, 300, 305, 316, 317, 343, 344, 350, 352, 354, 374, 376, 392, 407, 411-413, 429-432, 434, 435, 443, 474, 475, 498-501, 518, 687, 692, 693, 804, 862, 883, 884, 888, 890, and 891.

The immunostimulatory nucleic acids can be double-stranded or single-stranded.

5 Generally, double-stranded molecules are more stable *in vivo*, while single-stranded molecules have increased immune activity. Thus in some aspects of the invention it is preferred that the nucleic acid be single stranded and in other aspects it is preferred that the nucleic acid be double stranded.

The term T-rich nucleic acid and TG nucleic acid, as used herein, refers to an  
10 immunostimulatory T-rich nucleic acid and an immunostimulatory TG nucleic acid, respectively, unless otherwise indicated. The T-rich nucleic acid sequences of the invention are those broadly described above as well as the nucleic acids shown in Table A that have at least one poly T motif and/or have a composition of greater than 25% T or preferably 35% nucleotide residues. The C-rich nucleic acids are those having at least  
15 one and preferably at least two poly-C regions. The TG nucleic acids of the invention are those broadly described above as well as the specific nucleic acids shown in Table A that have at least one TG motif.

The nucleic acids of the invention may, but need not, also include a poly G motif. Poly G containing nucleic acids are also immunostimulatory. A variety of references,  
20 including Pisetsky and Reich, 1993 *Mol. Biol. Reports*, 18:217-221; Krieger and Herz, 1994, *Ann. Rev. Biochem.*, 63:601-637; Macaya et al., 1993, *PNAS*, 90:3745-3749; Wyatt et al., 1994, *PNAS*, 91:1356-1360; Rando and Hogan, 1998, In *Applied Antisense Oligonucleotide Technology*, ed. Krieg and Stein, p. 335-352; and Kimura et al., 1994, *J. Biochem.* 116, 991-994 also describe the immunostimulatory properties of poly G  
25 nucleic acids.

Poly G nucleic acids preferably are nucleic acids having the following formulas:



wherein  $X_1$ ,  $X_2$ ,  $X_3$ , and  $X_4$  are nucleotides. In preferred embodiments at least one of  $X_3$  and  $X_4$  are a G. In other embodiments both of  $X_3$  and  $X_4$  are a G. In yet other  
30 embodiments the preferred formula is 5' GGGN GGG 3', or 5' GGGN GGGN GGG 3' wherein N represents between 0 and 20 nucleotides. In other embodiments the poly G nucleic acid is free of unmethylated CG dinucleotides, such as, for example, the nucleic

- 34 -

acids listed below as SEQ ID NO: 5, 6, 73, 215, 267-269, 276, 282, 288, 297-299, 355, 359, 386, 387, 444, 476, 531, 557-559, 733, 768, 795, 796, 914-925, 928-931, 933-936, and 938. In other embodiments the poly G nucleic acid includes at least one unmethylated CG dinucleotide, such as, for example, the nucleic acids listed above as

5 SEQ ID NO: 67, 80-82, 141, 147, 148, 173, 178, 183, 185, 214, 224, 264, 265, 315, 329, 434, 435, 475, 519, 521-524, 526, 527, 535, 554, 565, 609, 628, 660, 661, 662, 725, 767, 825, 856, 857, 876, 892, 909, 926, 927, 932, and 937.

The terms "nucleic acid" and "oligonucleotide" are used interchangeably to mean multiple nucleotides (i.e. molecules comprising a sugar (e.g. ribose or deoxyribose) linked to a phosphate group and to an exchangeable organic base, which is either a

10 substituted pyrimidine (e.g. cytosine (C), thymidine (T) or uracil (U)) or a substituted purine (e.g. adenine (A) or guanine (G)). As used herein, the terms refer to oligoribonucleotides as well as oligodeoxyribonucleotides. The terms shall also include polynucleosides (i.e. a polynucleotide minus the phosphate) and any other organic base

15 containing polymer. Nucleic acid molecules can be obtained from existing nucleic acid sources (e.g., genomic or cDNA), but are preferably synthetic (e.g. produced by nucleic acid synthesis).

The terms nucleic acid and oligonucleotide also encompass nucleic acids or oligonucleotides with substitutions or modifications, such as in the bases and/or sugars.

20 For example, they include nucleic acids having backbone sugars which are covalently attached to low molecular weight organic groups other than a hydroxyl group at the 3' position and other than a phosphate group at the 5' position. Thus modified nucleic acids may include a 2'-O-alkylated ribose group. In addition, modified nucleic acids may include sugars such as arabinose instead of ribose. Thus the nucleic acids may be

25 heterogeneous in backbone composition thereby containing any possible combination of polymer units linked together such as peptide- nucleic acids (which have amino acid backbone with nucleic acid bases). In some embodiments, the nucleic acids are homogeneous in backbone composition. Nucleic acids also include substituted purines and pyrimidines such as C-5 propyne modified bases (Wagner et al., *Nature*

30 *Biotechnology* 14:840- 844, 1996). Purines and pyrimidines include but are not limited to adenine, cytosine, guanine, thymidine, 5-methylcytosine, 2-aminopurine, 2-amino-6-chloropurine, 2,6-diaminopurine, hypoxanthine, and other naturally and



non-naturally occurring nucleobases, substituted and unsubstituted aromatic moieties. Other such modifications are well known to those of skill in the art.

For use in the instant invention, the nucleic acids of the invention can be synthesized *de novo* using any of a number of procedures well known in the art. For example, the b-cyanoethyl phosphoramidite method (Beaucage, S.L., and Caruthers, M.H., *Tet. Let.* 22:1859, 1981); nucleoside H-phosphonate method (Garegg *et al.*, *Tet. Let.* 27:4051-4054, 1986; Froehler *et al.*, *Nucl. Acid. Res.* 14:5399-5407, 1986, ; Garegg *et al.*, *Tet. Let.* 27:4055-4058, 1986, Gaffney *et al.*, *Tet. Let.* 29:2619-2622, 1988).

These chemistries can be performed by a variety of automated nucleic acid synthesizers available in the market. These nucleic acids are referred to as synthetic nucleic acids. Alternatively, T-rich and/or TG dinucleotides can be produced on a large scale in plasmids, (see Sambrook, T., *et al.*, "Molecular Cloning: A Laboratory Manual", Cold Spring Harbor laboratory Press, New York, 1989) and separated into smaller pieces or administered whole. Nucleic acids can be prepared from existing nucleic acid sequences (e.g., genomic or cDNA) using known techniques, such as those employing restriction enzymes, exonucleases or endonucleases. Nucleic acids prepared in this manner are referred to as isolated nucleic acid. An isolated nucleic acid generally refers to a nucleic acid which is separated from components which it is normally associated with in nature. As an example, an isolated nucleic acid may be one which is separated from a cell, from a nucleus, from mitochondria or from chromatin. The terms Py-rich nucleic acids and TG nucleic acids encompasses both synthetic and isolated Py-rich nucleic acids and TG nucleic acids.

For use *in vivo*, the Py-rich and TG nucleic acids may optionally be relatively resistant to degradation (e.g., are stabilized). A "stabilized nucleic acid molecule" shall mean a nucleic acid molecule that is relatively resistant to *in vivo* degradation (e.g. via an exo- or endo-nuclease). Stabilization can be a function of length or secondary structure. Nucleic acids that are tens to hundreds of kbs long are relatively resistant to *in vivo* degradation. For shorter nucleic acids, secondary structure can stabilize and increase their effect. For example, if the 3' end of an nucleic acid has self-complementarity to an upstream region, so that it can fold back and form a sort of stem loop structure, then the nucleic acid becomes stabilized and therefore exhibits more activity.

- 36 -

Alternatively, nucleic acid stabilization can be accomplished via phosphate backbone modifications. Preferred stabilized nucleic acids of the instant invention have a modified backbone. It has been demonstrated that modification of the nucleic acid backbone provides enhanced activity of the Py-rich and TG nucleic acids when administered *in vivo*. These stabilized structures are preferred because the Py-rich and TG molecules of the invention have at least a partial modified backbone. Py-rich and TG constructs having phosphorothioate linkages provide maximal activity and protect the nucleic acid from degradation by intracellular exo- and endo-nucleases. Other modified nucleic acids include phosphodiester modified nucleic acids, combinations of phosphodiester and phosphorothioate nucleic acid, methylphosphonate, methylphosphorothioate, phosphorodithioate, p-ethoxy, and combinations thereof. Each of these combinations and their particular effects on immune cells is discussed in more detail with respect to CpG nucleic acids in PCT Published Patent Applications PCT/US95/01570 (WO 96/02555) and PCT/US97/19791 (WO 98/18810) claiming priority to U.S. Serial Nos. 08/386,063 and 08/960,774, filed on February 7, 1995 and October 30, 1997 respectively, the entire contents of which are hereby incorporated by reference. It is believed that these modified nucleic acids may show more stimulatory activity due to enhanced nuclease resistance, increased cellular uptake, increased protein binding, and/or altered intracellular localization.

The compositions of the invention may optionally be chimeric oligonucleotides. The chimeric oligonucleotides are oligonucleotides having a formula: 5' Y<sub>1</sub>N<sub>1</sub>ZN<sub>2</sub>Y<sub>2</sub> 3'. Y<sub>1</sub> and Y<sub>2</sub> are nucleic acid molecules having between 1 and 10 nucleotides. Y<sub>1</sub> and Y<sub>2</sub> each include at least one modified internucleotide linkage. Since at least 2 nucleotides of the chimeric oligonucleotides include backbone modifications these nucleic acids are an example of one type of "stabilized immunostimulatory nucleic acids."

With respect to the chimeric oligonucleotides, Y<sub>1</sub> and Y<sub>2</sub> are considered independent of one another. This means that each of Y<sub>1</sub> and Y<sub>2</sub> may or may not have different sequences and different backbone linkages from one another in the same molecule. The sequences vary, but in some cases Y<sub>1</sub> and Y<sub>2</sub> have a poly-G sequence. A poly-G sequence refers to at least 3 Gs in a row. In other embodiments the poly-G sequence refers to at least 4, 5, 6, 7, or 8 Gs in a row. In other embodiments Y<sub>1</sub> and Y<sub>2</sub> may be TCGTCG, TCGTCGT, or TCGTCGTT (SEQ ID NO:1145). Y<sub>1</sub> and Y<sub>2</sub> may also

- 37 -

have a poly-C, poly-T, or poly-A sequence. In some embodiments  $Y_1$  and/or  $Y_2$  have between 3 and 8 nucleotides.

$N_1$  and  $N_2$  are nucleic acid molecules having between 0 and 5 nucleotides as long as  $N_1ZN_2$  has at least 6 nucleotides in total. The nucleotides of  $N_1ZN_2$  have a

5 phosphodiester backbone and do not include nucleic acids having a modified backbone.

Z is an immunostimulatory nucleic acid motif but does not include a CG. For instance, Z may be a nucleic acid a T-rich sequence, e.g. including a TTTT motif or a sequence wherein at least 50% of the bases of the sequence are Ts or Z may be a TG sequence.

10 The center nucleotides ( $N_1ZN_2$ ) of the formula  $Y_1N_1ZN_2Y_2$  have phosphodiester internucleotide linkages and  $Y_1$  and  $Y_2$  have at least one, but may have more than one or even may have all modified internucleotide linkages. In preferred embodiments  $Y_1$  and/or  $Y_2$  have at least two or between two and five modified internucleotide linkages or  $Y_1$  has two modified internucleotide linkages and  $Y_2$  has five modified internucleotide  
15 linkages or  $Y_1$  has five modified internucleotide linkages and  $Y_2$  has two modified internucleotide linkages. The modified internucleotide linkage, in some embodiments is a phosphorothioate modified linkage, a phosphorodithioate modified linkage or a p-ethoxy modified linkage.

Modified backbones such as phosphorothioates may be synthesized using  
20 automated techniques employing either phosphoramidate or H-phosphonate chemistries. Aryl- and alkyl-phosphonates can be made, e.g., as described in U.S. Patent No. 4,469,863; and alkylphosphotriesters (in which the charged oxygen moiety is alkylated as described in U.S. Patent No. 5,023,243 and European Patent No. 092,574) can be prepared by automated solid phase synthesis using commercially available reagents.  
25 Methods for making other DNA backbone modifications and substitutions have been described (Uhlmann, E. and Peyman, A., *Chem. Rev.* 90:544, 1990; Goodchild, J., *Bioconjugate Chem.* 1:165, 1990).

Other stabilized nucleic acids include: nonionic DNA analogs, such as alkyl- and aryl-phosphates (in which the charged phosphonate oxygen is replaced by an alkyl or  
30 aryl group), phosphodiester and alkylphosphotriesters, in which the charged oxygen moiety is alkylated. Nucleic acids which contain diol, such as tetraethyleneglycol or

- 38 -

hexaethyleneglycol, at either or both termini have also been shown to be substantially resistant to nuclease degradation.

In the case when the Py-rich or TG nucleic acid is administered in conjunction with an antigen which is encoded in a nucleic acid vector, it is preferred that the backbone of the Py-rich or TG nucleic acid be a chimeric combination of phosphodiester and phosphorothioate (or other phosphate modification). The cell may have a problem taking up a plasmid vector in the presence of completely phosphorothioate nucleic acid. Thus when both a vector and a nucleic acid are delivered to a subject, it is preferred that the nucleic acid have a chimeric backbone or have a phosphorothioate backbone but that the plasmid be associated with a vehicle that delivers it directly into the cell, thus avoiding the need for cellular uptake. Such vehicles are known in the art and include, for example, liposomes and gene guns.

The nucleic acids described herein as well as various control nucleic acids are presented below in Table A.

Table A

SEQ ID NO:	ODN SEQUENCE	BACKBONE
1	tctcccagcgtgcgccat	s
2	ataatccagcttgaaccaag	s
3	ataatcgacgttcaagcaag	s
4	taccgcgtgcgaccctct	s
5	ggggaggggt	s
6	ggggagggg	s
7	ggtgaggtg	s
8	tccatgtzgttcctgatgct	o
9	gctaccttagzgtga	o
10	tccatgazgttcctgatgct	o
11	tccatgacgttcztgatgct	o
12	gctagazgttagtgt	o
13	agctccatggtgctcactg	s
14	ccacgtcgaccctcaggcga	s
15	gcacatcgtcccgcagccga	s
16	gtcactcgtggtacctcga	s
17	gttggatacaggccagactttgttg	o
18	gattcaacttgcgctcatcttaggc	o
19	accatggacgaactgtttcccctc	s
20	accatggacgagctgtttcccctc	s
21	accatggacgacctgtttcccctc	s
22	accatggacgtactgtttcccctc	s
23	accatggacggtctgtttcccctc	s
24	accatggacgttctgtttcccctc	s
25	ccactcacatctgctgtccacaag	o
26	acttctcatagtccctttggtccag	o

- 39 -

27	tccatgagcttcctgagtct	o
28	gaggaaggigiggaigacgt	o
29	gtgaaticggttcicgggict	o
30	aaaaaa	s
31	cccccc	s
32	ctgtca	s
33	tcgtag	s
34	tcgtgg	s
35	cgtcgt	s
36	tccatgtcggctcctgagtct	sos
37	tccatgccggctcctgagtct	sos
38	tccatgacggctcctgagtct	sos
39	tccatgacggctcctgagtct	sos
40	tccatgtcgatcctgagtct	sos
41	tccatgtcgctcctgagtct	sos
42	tccatgtcgttcctgagtct	sos
43	tccatgacgttcctgagtct	sos
44	tccataacgttcctgagtct	sos
45	tccatgacgtccctgagtct	sos
46	tccatcacgtgcctgagtct	sos
47	tccatgctggctcctgagtct	sos
48	tccatgtzggctcctgagtct	sos
49	cgcgttcctccagatgagctcatgggtttctccaccaag	o
50	cttgggtggagaaacccatgagctcatctggaggaagcgg	o
51	cccacaaaggatgagaagtt	o
52	agatagcaaatacggtgacg	o
53	ggttcacgtgctcatggctg	o
54	tctcccagcgtgcgccat	s
55	tctcccagcgtgcgccat	s
56	taccgcgtgcgacctct	s
57	ataatccagcttgaaccaag	s
58	ataatcgacgttcaagcaag	s
59	tccatgattttcctgatttt	o
60	ttgtttttttgtttttttgttttt	s
61	ttttttttgtttttttgttttt	o
62	tgtgtctttgtgtctttgtgtct	s
63	tgtgtctgtgtctttgtgtct	o
64	gcattcatcaggcgggcaagaat	o
65	taccgagcttcgacgagatttca	o
66	gcatgacgttgagct	s
67	cacgttgaggggcat	s
68	ctgctgagactggag	s
69	tccatgacgttcctgacgtt	s
70	gcatgagcttgagctga	o
71	tcagcgtgcgcc	s
72	atgacgttcctgacgtt	s
73	ttttggggttttggggtttt	s
74	tctaggctttttaggcttcc	s
75	tgcattttttaggccaccat	s
76	tctcccagcgtgcgtgcgccat	s
77	tctcccagcgggcgcat	s

78	tctcccagcgagcgccat	s
79	tctcccagcgcgccat	s
80	ggggtgacgttcagggggg	sos
81	ggggtccagcgtgcgccatggggg	sos
82	ggggtgtcgttcagggggg	sos
83	tccatgtcgttcctgtcgtt	s
84	tccatagcgttcctagcgtt	s
85	tcgtcgtcgttcctcgttctt	s
86	gcatgacgttgagct	sos
87	tctcccagcgtgcgccatat	sos
88	tccatgazgttcctgazgtt	s
89	gcatgazgttgagct	o
90	tccagcgtgcgccata	sos
91	tctcccagcgtgcgccat	o
92	tccatgagcttcctgagtct	o
93	gcatgtcgttgagct	sos
94	tcctgacgttcctgacgtt	s
95	gcatgatgttgagct	o
96	gcatttcgaggagct	o
97	gcatgtagctgagct	o
98	tccaggacgttcctagttct	o
99	tccaggagcttcctagttct	o
100	tccaggatgttcctagttct	o
101	tccagtctaggcctagttct	o
102	tccagttcgagcctagttct	o
103	gcatggcgttgagct	sos
104	gcatagcgttgagct	sos
105	gcattgcgttgagct	sos
106	gcttgcgttgcgttt	sos
107	tctcccagcgttcgccatat	sos
108	tctcccagcgtgcgttatat	sos
109	tctccctgcgtgcgccatat	sos
110	tctgcgtgcgtgcgccatat	sos
111	tctcctagcgtgcgccatat	sos
112	tctcccagcgtgcgcctttt	sos
113	gctandcghhagc	o
114	tcctgacgttccc	o
115	ggaagacgttaga	o
116	tcctgacgttaga	o
117	tcagaccagctgggtcggtgttcctga	o
118	tcaggaacacccgaccagctgggtctga	o
119	gctagtcgatagc	o
120	gctagtcgctagc	o
121	gcttgacgtctagc	o
122	gcttgacgttttagc	o
123	gcttgacgtcaagc	o
124	gctagacgttttagc	o
125	tccatgacattcctgatgct	o
126	gctagacgtctagc	o
127	ggctatgtcgttcctagcc	o
128	ggctatgtcgatcctagcc	o

- 41 -

129	ctcatggggtttctccaccaag	o
130	cttgggtggagaaacccatgag	o
131	tccatgacgttcctagttct	o
132	ccgcttcctccagatgagctcatg	o
133	catgagctcatctggaggaagcgg	o
134	ccagatgagctcatgggtttctcc	o
135	ggagaaacccatgagctcatctgg	o
136	agcatcaggaacgacatgga	o
137	tccatgacgttcctgacgtt	rna
138	gcgcgcgcgcgcgcgcgcgcg	o
139	ccggccggccggccggccgg	o
140	ttccaatcagccccacccgctctggccccaccctcaccctcca	o
141	tggaggtgaggggtggggccagagcgggtggggctgattggaa	o
142	tcaaagtgggattttcccatgagtct	o
143	agactcatgggaaaatcccacatttga	o
144	tccaagtgcctgagtcactaataaaga	o
145	tctttattagtgcactcagcacttgga	o
146	tgcaggaagtcgggttttccccaaaccccc	o
147	ggggggttggggaaaacccggacttctgca	o
148	ggggactttccgctgggactttccagggggactttcc	sos
149	tccatgacgttcctctccatgacgttcctctccatgacgttcctc	o
150	gaggaacgtcatggagaggaacgtcatggagaggaacgtcatgga	o
151	ataatagagcttcaagcaag	s
152	tccatgacgttcctgacgtt	s
153	tccatgacgttcctgacgtt	sos
154	tccaggactttcctcagggtt	s
155	tcttgcatgctaaaggacgtcacattgcacaatcttaataaggt	o
156	accttattaagattgtgcaatgtgacgtccttttagcatcgcaaga	o
157	tcctgacgttcctggcggtcctgtcgct	o
158	tcctgtcgctcctgtcgct	o
159	tcctgacgttgaagt	o
160	tcctgtcggtgaagt	o
161	tcctggcggtgaagt	o
162	tcctgccgttgaagt	o
163	tccttacgttgaagt	o
164	tcctaacgttgaagt	o
165	tcctcacgttgaagt	o
166	tcctgacgatgaagt	o
167	tcctgacgtgaagt	o
168	tcctgacggtgaagt	o
169	tcctgacgtagaagt	o
170	tcctgacgtcgaagt	o
171	tcctgacgtggaagt	o
172	tcctgagcttgaagt	o
173	gggggacgttggggg	o
174	tcctgacgttccttc	o
175	tctcccagcgagcgagcgccat	s
176	tcctgacgttccccctggcggtccccctgtcgct	o
177	tcctgtcgctcctgtcgctcctgtcgct	o
178	tcctggcggggaagt	o
179	tcctgazgttgaagt	o

- 42 -

180	tcztgacgttgaagt	o
181	tcctagcgttgaagt	o
182	tccagacgttgaagt	o
183	tcctgacggggaagt	o
184	tcctggcgtgaagt	o
185	ggctccggggagggaattttgtctat	o
186	atagacaaaaattccctccccggagcc	o
187	tccatgagcttccttgagtct	rna
188	tcgtcgtctctccgcttctt	so
189	tcgtcgtctctccgcttctt	s20
190	tcgagacattgcacaatcatctg	o
191	cagattgtgcaatgtctcga	o
192	tccatgtcgttcctgatgcg	o
193	gcgatgtcgttcctgatgct	o
194	gcgatgtcgttcctgatgcg	o
195	tccatgtcgttcgcgcgcg	o
196	tccatgtcgttcctgccgct	o
197	tccatgtcgttcctgtagct	o
198	gcggcgggcgcgcgcgcc	o
199	atcaggaacgtcatgggaagc	o
200	tccatgagcttcctgagtct	p-ethoxy
201	tcaacgtt	p-ethoxy
202	tcaagctt	p-ethoxy
203	tcctgtcgttcctgtcgtt	s
204	tccatgtcgtttttgtcgtt	s
205	tcctgtcgttcctgtcgtt	s
206	tcctgtcgttcctgtcgtt	s
207	btccattccatgacgttcctgatgcttcca	os
208	tcctgtcgtttttgtcgtt	s
209	tcgtcgtctctccgcttctt	s
210	tcgtcgtctctgcccttctt	s
211	tcgtcgtctgtgtcgttctt	s
212	tcctgtcgttcctgtcgttgaacgacagg	o
213	tcctgtcgttcctgtcgttcaacgtcaggaacgacagga	o
214	ggggctctgtcgttttggggg	sos
215	ggggctctgtcgttttggggg	sos
216	tccggccgttgaagt	o
217	tccggacggtgaagt	o
218	tcccggcgttgaagt	o
219	tccagacggtgaagt	o
220	tcccagcgttgaagt	o
221	tccagagcttgaagt	o
222	tccatgtzgttcctgtzgtt	s
223	tccatgacgttcctgacgtt	sos
224	gggggtgacgttttggggg	sos
225	tccaggacttctctcaggtt	s
226	tttttttttttttttttt	s
227	tccatgccgttcctgccgtt	s
228	tccatggcgggectggcggg	s
229	tccatgacgttcctgccgtt	s
230	tccatgacgttcctggcggg	s



231	tccatgacgttcctgcgttt	s
232	tccatgacggtcctgacggt	s
233	tccatgcgtgcgtgcgtttt	s
234	tccatgcgttgcggtgcgtt	s
235	btccattccattctaggcctgagtccttccat	os
236	tccatagcgttcctagcgtt	o
237	tccatgtcggttcctgtcggt	o
238	tccatagcgatcctagcgat	o
239	tccattgcgttccttgcgtt	o
240	tccatagcggtcctagcggg	o
241	tccatgattttcctgcagttcctgatttt	
242	tccatgacgttcctgcagttcctgacggt	s
243	ggcggcgggcgggcgggcg	o
244	tccacgacgttttcgacggt	s
245	tcgtcggttgctggttgctggt	s
246	tcgtcggtttgtcggtttgtcggt	s
247	tcgtcggttgctggtttgtcggt	s
248	gcgtgcgttgctggttgctggt	s
249	czggczggczggczccgg	o
250	gcggcgggcgggcgcgcccc	s
251	agicccgigaacgiattcac	o
252	tgctggttgctggttgctggt	s
253	tgctggttgctggttgctggt	s
254	tgctggttgctggttgctggt	s
255	tcgtcggtcgctggt	s
256	tgctggttgctggt	s
257	cccccccccccccccccc	s
258	tctagcgttttttagcgttcc	sos
259	tgcattccccagggccaccat	s
260	tcgtcggtcgctgctgctgctggt	sos
261	tcgtcggttgctggttgctggt	sos
262	tcgtcggtttgtcggtttgtcggt	sos
263	tcgtcggttgctggtttgtcggt	sos
264	ggggaggaggaaacttcttaaaattccccccagaatggtt	o
265	aaacattctgggggaattttaagaagttcctccctcccc	o
266	atgtttacttcttaaaattccccccagaatggtt	o
267	aaacattctgggggaattttaagaagtaaacat	o
268	atgtttactagacaaaattccccccagaatggtt	o
269	aaacattctgggggaattttgtctagtaaacat	o
270	aaaattgacgtttttaaaaa	sos
271	ccccttgacgttttcccccc	sos
272	tttctggttggtttgtcggt	
273	tcgtcggtttgtcggtttgtcggt	sos
274	ctgcagcctgggac	o
275	acccgtcgtaattatagtaaaaccc	o
276	ggtacctgtggggacattgtg	o
277	agcaccgaacgtgagagg	o
278	tccatgccgttcctgccgtt	o
279	tccatgacggtcctgacggt	o
280	tccatgccggtcctgccggt	o
281	tccatgcgcgtcctgcgcgt	o

- 44 -

282	ctggtctttctggttttttctgg	s
283	tcaggggtggggggaacctt	sos
284	tccatgazgttcctagttct	o
285	tccatgatgttcctagttct	o
286	cccgaagtcatttcctcttaacctgg	o
287	ccaggttaagaggaaatgacttcggg	o
288	tcctggzggggaagt	o
289	gzggzgggzzggzgzgcccc	x
290	tccatgtgcttcctgatgct	o
291	tccatgtccttcctgatgct	
292	tccatgtcgttcctagttct	
293	tccaagtagttcctagttct	o
294	tccatgtagttcctagttct	o
295	tcgcgcgcgttcgcgcggtt	s
296	tcctggcggctcctggcggtt	s
297	tcctggaggggaagt	o
298	tcctgggggggaagt	o
299	tcctgggtggggaagt	o
300	tcgtcgttttgtcgttttgtcggt	o
301	ctggtctttctggttttttctgg	o
302	tccatgacgttcctgacgtt	o
303	tccaggacttctctcaggtt	sos
304	tzgtzgttttgtzgttttztggt	o
305	btcgtcgttttgtcgttttgtcggtttttt	os
306	gctatgacgttccaagg	s
307	tcaacgtt	s
308	tccaggactttcctcaggtt	o
309	ctctctgtaggcccgcttgg	s
310	ctttccgttggaccctggg	s
311	gtccggggccaggccaaagtc	s
312	gtgcgcgcgagcccgaaatc	s
313	tccatgaigtctcctgaigt	s
314	aatagtcgcataacaaaac	o
315	aatagtcgcatggcggggc	o
316	btttttccatgtcggtcctgatgcttttt	os
317	tcctgtcggtgaagttttt	o
318	gctagcttttagagcttttagagctt	o
319	tgctgcttcccccccccc	o
320	tcgacgttcccccccccc	o
321	tcgtcggtcccccccccc	o
322	tcgtcggtcccccccccc	o
323	tcgccgttcccccccccc	o
324	tcgtcgatcccccccccc	o
325	tcctgacgttgaagt	s
326	tcctgccgttgaagt	s
327	tcctgacggtgaagt	s
328	tcctgagcttgaagt	s
329	tcctggcggggaagt	s
330	aaaatctgtgcttttaaaaaa	sos
331	gatccagtcacagtgaacctggcagaatctggat	o
332	gatccagattctgccaggtcactgtgactggat	o

- 45 -

333	gatccagtcacagtgactcagcagaatctggat	o
334	gatccagattctgctgagtcactgtgactggat	o
335	tcgtcggtccccccczcccc	o
336	tzgtqggtcccccccccccc	o
337	tzgtcggtcccccccccccc	o
338	tcgtzgttcccccccccccc	o
339	tcgtcggtcccccccccccc	o
340	tcgtcggtcccccccccccc	o
341	tcggcggtcccccccccccc	o
342	ggccttttcccccccccccc	o
343	tcgtcggttttgacgttttgctggt	s
344	tcgtcggttttgacgttttgacgtt	s
345	ccgtcggtcccccccccccc	o
346	gcgtcggtcccccccccccc	o
347	tcgtcattcccccccccccc	o
348	acgtcggtcccccccccccc	o
349	ctgtcggtcccccccccccc	o
350	btttttcgtcggtcccccccccccc	os
351	tcgtcggtcccccccccccc	o
352	tcgtcggttttgctggttttgctggtb	o
353	tccagttccttcctcagttc	o
354	tzgtcggttttgctggttttgctggt	o
355	tcctggaggggaagt	s
356	tcctgaaaaggaagt	s
357	tcgtcggtcccccccc	s
358	tzgtzgttttgztggtttgztggt	s
359	ggggtcaagcttgagggggg	sos
360	tgctgcttcccccccccccc	s
361	tcgtcggtcggt	s2
362	tcgtcggtcggt	s20
363	tcgtcggtcggt	os2
364	tcaacgttga	s
365	tcaacgtt	s
366	atagttttccatttttttac	
367	aatagtcgcatcgcgac	o
368	aatagtcgcatcccgac	o
369	aatagtcgcatcccccc	o
370	tgctgcttttgctgttttgctgt	o
371	ctgtgctttctgtgttttctgtg	s
372	ctaacttttctaatttttttctaa	s
373	tcgtcggtggtgctggtgctggt	s
374	tcgtcggtggtgctggtttggt	s
375	accatggacgagctgttcccctc	
376	tcgtcggtttgctgctgtt	s
377	ctgtaagtgagcttgagag	
378	gagaacgctggaccttcc	
379	cgggcgactcagttctatcgg	
380	gttctcagataaagcggaaccagcaacagacacagaa	
381	ttctgtgtctgttgctggttccgctttatctgagaac	
382	cagacacagaagcccgatagacg	
383	agacagacacgaaacgaccg	

- 46 -

384	gtctgtcccatgatctcgaa	
385	gctggccagcttacctccc	
386	ggggcctctatacaacctggg	
387	ggggccctgagactgcc	
388	gagaacgctggaccttccat	
389	tccatgtcggtcctgatgct	
390	ctcttgcgacctggaaggta	
391	aggtacagccaggactacga	
392	accatggacgacctgtttcccctc	
393	accatggattacctttttcccctt	
394	atggaaggtccagcggtctc	o
395	agcatcaggaccgacatgga	o
396	ctctccaagctcacttacag	
397	tccctgagactgccccacctt	
398	gccacaaaacttgccatg	
399	gtccatggcgtgcgggatga	
400	cctctatacaacctgggac	
401	cgggcgactcagtcctatcgg	
402	gcgctaccggtagcctgagt	
403	cgactgccgaacaggatatcggtgatcagcactgg	
404	ccagtgcgtgatcaccgatatcctgttcggcagtcg	
405	ccaggttgtatagaggc	
406	tctcccagcgtacgccat	s
407	tctcccagcgtgcgtttt	s
408	tctcccagcgtgcgccat	s
409	tctcccgtcgtgcgccat	s
410	ataatcgtcgttcaagcaag	s
411	tcgtcgttttgtcgttttgtcggt	s2
412	tcgtcgttttgtcgttttgtcggt	s2
413	tcgtcgttttgtcgttttgtcggt	s2
414	tcttcgtnttntcgtnttntcgtnt	s
415	tctcccagcgtcgccat	s
416	tctcccacgtcgccat	s
417	ataatcgtcgttcaagaaag	s
418	ataatcgacgttcccccccc	s
419	tctatcgacgttcaagcaag	s
420	tcc tga cgg gg agt	s
421	tccatgacgttcctgatcc	
422	tccatgacgttcctgatcc	
423	tccatgacgttcctgatcc	
424	tcc tgg cgt gga agt	s
425	tccatgacgttcctgatcc	
426	tcgtcgctgttgctggtttctt	s
427	agcagcttttagagcttttagagctt	s
428	cccccccccccccccccccc	s
429	tcgtcgttttgtcgttttgtcgttttgtcggt	s
430	tcgtcgttttttgtcgttttttgtcggt	s
431	tcgtcgttttttttttttttt	s
432	tttttcaacggttgatttttt	sos
433	ttttttttttttttttttttt	s
434	ggggtcgtcgttttgggggg	

- 47 -

435	tcgtcggtttgtcggtttggggg	
436	tcgtcgctgtctccgcttcttcttgcc	s
437	tcgtcgctgtctccg	s
438	ctgtaagtgaagcttgagag	
439	gagaacgctggaccttccat	
440	ccagggtgtatagaggc	
441	gctagacgttagcgtga	
442	ggagctcttcgaacgccata	
443	tctccatgatggtttatcg	
444	aaggtggggcagtctcagga	
445	atcggaggactggcgccg	
446	ttaggacaaggtctagggtg	
447	accacaacgagaggaacgca	
448	ggcagtgcaggctcaccggg	
449	gaaccttccatgctgtt	
450	gctagacgttagcgtga	
451	gcttggagggcctgtaagtg	
452	gtagccttccta	
453	cggtagccttccta	
454	cacggtagccttccta	
455	agcacggtagccttccta	
456	gaacgctggaccttccat	
457	gaccttccat	
458	tggaccttccat	
459	gctggaccttccat	
460	acgctggaccttccat	
461	taagctctgtcaacgccagg	
462	gagaacgctggaccttccatgt	
463	tccatgtcggtcctgatgct	
464	ttcatgccttgcaaaatggcg	
465	tgctagctgtgcctgtacct	
466	agcatcaggaccgacatgga	
467	gaccttccatgtcggtcctgat	
468	acaaccacgagaacgggaac	
469	gaaccttccatgctgttccg	
470	caatcaatctgaggagaccc	
471	tcagctctggtacttttca	
472	tggttacggtctgtcccatg	
473	gtctatcggaggactggcg	
474	cattttacgggcggcgggc	
475	gaggggaccattttacgggc	
476	tgtccagccgaggggacat	
477	cgggcttacggcggtgctg	
478	tggaccttctatgtcggtcc	
479	tgtcccatgtttttagaagc	
480	gtggttacgggtcggtcccat	
481	cctccaaatgaaagaccccc	
482	ttgtactctccatgatggtt	
483	ttccatgctgttccggctgg	
484	gaccttctatgtcggtcctg	
485	gagaccgctcgaccttcgat	

486	ttgccccatatttttagaaac	
487	ttgaaactgaggtgggac	
488	ctatcggaggactggcgccg	
489	cttgaggggcctcccggcg	
490	gctgaaccttccatgctgtt	
491	tagaaacagcattcttcttttagggcagcaca	
492	agatggttctcagataaagcgaa	
493	ttccgctttatctgagaaccatct	
494	gtcccaggttgatatagaggctgc	
495	gcgccagtcctccgatagac	
496	atcggaggactggcgcccg	
497	ggtctgtcccatatttttag	
498	tttttcaacggttgagggggg	sos
499	tttttcaacggttgatttttt	sos
500	ggggtcaacggttgatttttt	sos
501	ggggttttcaacggtttgagggggg	sos
502	ggttacggtctgtcccatat	
503	ctgtcccatatttttagaca	
504	accatcctgaggccattcgg	
505	cgtctatcgggcttctgtgtctg	
506	ggccatcccacattgaaagt	
507	ccaaatatcgggtgtcaagcac	
508	gtgcttgaccaccgatatttgg	
509	gtgctgatcaccgatatcctgttcgg	
510	ggccaactttcaatgtgggatggcctc	
511	ttccgccgaatggcctcaggatggtac	
512	tatagtccctgagactgccccaccttctcaacaacc	
513	gcagcctctatacaacctgggacggga	
514	ctatcggaggactggcgcccg	
515	tatcggaggactggcgcccg	
516	gatcggaggactggcgcccg	
517	ccgaacaggatatcgggtgatcagcac	
518	ttttgggtcaacggttgagggggg	
519	ggggtcaacggttgagggggg	sos
520	cgcgcgcgcgcgcgcgcgcg	s
521	ggggcatgacgttcggggggg	ss
522	ggggcatgacgttcaaaaaa	s
523	ggggcatgagcttcggggggg	s
524	ggggcatgacgttcggggggg	sos
525	aaaacatgacgttcaaaaaa	sos
526	aaaacatgacgttcggggggg	sos
527	ggggcatgacgttcaaaaaa	sos
528	accatggacgatctgtttccctc	s
529	gccatggacgaactgtttccctc	s
530	cccccccccccccccccc	sos
531	gggggggggggggggggggg	sos
532	gctgtaaaatgaatcggccg	sos
533	ttcgggaggactcctccatt	sos
534	tatgccgcgcccggacttat	sos
535	ggggtaatcgatcagggggg	sos
536	tttgagaacgctggaccttc	sos

- 49 -

537	gatcgctgatctaatactcg	sos
538	gtcggtcctgatgctgttcc	sos
539	tcgtcgtcagttcgctgtcg	sos
540	ctggaccttccatgtcgg	sos
541	gctcgttcagcgcgtct	sos
542	ctggaccttccatgtc	sos
543	caactgtccttcgtcga	sos
544	cgctggaccttccatgtcgg	sos
545	gctgagctcatgccgtctgc	sos
546	aacgctggaccttccatgtc	sos
547	tgcataccgtacacagctct	sos
548	ccttccatgtcggctctgat	sos
549	tactcttcggatcccttgcg	sos
550	ttccatgtcggctctgat	sos
551	ctgattgctctctcgtga	sos
552	ggcgttattcctgactcgcc	o
553	cctacgttgatgcgccagct	o
554	ggggtaatacgatgagggggg	o
555	ttcgggaggactcctccatt	o
556	tttttttttttttttttttt	o
557	gggggtttttttttttggggg	o
558	tttttgggggggggggttttt	o
559	gggggggggggggggggggt	o
560	aaaaaaaaaaaaaaaaaaaaa	o
561	cccccaaaaaaaaaaaccccc	o
562	aaaaaccccccccccaaaaa	o
563	tttgaattcaggactggtaggtgag	o
564	tttgaatcctcagcggctctccagtgc	o
565	aattctctatcgggcttctgtgtctgttgctgggtccgctttat	o
566	ctagataaagcggaaccagcaacagacacagaagccccgatagag	o
567	ttttctagagaggtgcacaatgctctgg	o
568	tttgaattccgtgtacagaagcgagaagc	o
569	tttgcggccgctagacttaacctgagagata	o
570	tttggggccacgagagacagagacacttc	o
571	tttggggccgcttctcgttctgtacacg	o
572	gagaacgctggaccttccat	s
573	tccatgtcggctcctgatgct	s
574	ctgtcg	s
575	tcgtga	s
576	cgtcga	s
577	agtgtc	s
578	ctgtcg	o
579	agtgtc	o
580	cgtcga	o
581	tcgtga	o
582	gagaacgctccagcttcgat	o
583	gctagacgtaagcgtga	o
584	gagaacgctcgaccttccat	o
585	gagaacgctggacctatccat	o
586	gctagaggtttagcgtga	o
587	gagaacgctggacttccat	o

- 50 -

588	tcacgctaacgtctagc	o
589	bgctagacggttagcgtga	o
590	atggaaggtcgagcgttctc	o
591	gagaacgctggaccttcgat	o
592	gagaacgatggaccttccat	o
593	gagaacgctggatccat	o
594	gagaacgctccagcactgat	o
595	tccatgtcgggtcctgctgat	o
596	atgtcctcgggtcctgatgct	o
597	gagaacgctccaccttccat	o
598	gagaacgctggaccttcgta	o
599	batggaaggtccagcgttctc	o
600	tcctga	o
601	tcaacggtt	o
602	aacggtt	o
603	aacggttga	o
604	tcacgctaacctctagc	o
605	gagaacgctggaccttgcgat	o
606	gctggaccttccat	o
607	gagaacgctggacctcatccat	o
608	gagaacgctggacgctcatccat	o
609	aacggttgaggggcat	o
610	atgccctcaacggtt	o
611	tcaacggttga	o
612	gctggaccttccat	o
613	caacggtt	o
614	acaacggttga	o
615	tcacgt	o
616	tcaagctt	o
617	tcgtca	o
618	aggatatc	o
619	tagacgtc	o
620	gacgtcat	o
621	ccatcgat	o
622	atcgatgt	o
623	atgcatgt	o
624	ccatgcat	o
625	agcgtga	o
626	tcagcgct	o
627	ccttcgat	o
628	gtgccggggtctccgggc	s
629	gctgtggggcggtcctg	s
630	btcaacggtt	o
631	ftcaacggtt	o
632	faacggttga	o
633	tcaacgt	s
634	aacggttg	s
635	cgacga	o
636	tcaacggtt	o
637	tcgga	o
638	agaacggtt	o



639	tcatcgat	o
640	taaacggt	s
641	ccaacggt	s
642	gctcga	s
643	cgacgt	s
644	cgtcgt	s
645	acgtgt	s
646	cgttcg	s
647	gagcaagctggaccttccat	s
648	cgcgta	s
649	cgtacg	s
650	tcaccggt	s
651	caagagatgctaacaatgca	s
652	acccatcaatagctctgtgc	s
653	ccatcgat	o
654	tcgacgtc	o
655	ctagcgct	o
656	taagcgct	o
657	tcgcgaattcgcg	o
658	atggaagggtccagcgttct	o
659	actggacgttagcgtga	o
660	cgcctggggctggtctgg	o
661	gtgtcggggtctccgggc	o
662	gtgccggggtctccgggc	o
663	cgcgctcgcggcggttgg	o
664	gaagttcacgttgaggggcat	o
665	atctggtgagggcaagctatg	s
666	gttgaaacccgagaacatcat	s
667	gcaacggt	o
668	gtaacggt	o
669	cgaacggt	o
670	gaaacggt	o
671	caaacggt	o
672	ctaacggt	o
673	ggaacggt	o
674	tgaacggt	o
675	acaacggt	o
676	ttaacggt	o
677	aaaacggt	o
678	ataacggt	o
679	aacgttct	o
680	tccgatcg	o
681	tccgtacg	o
682	gctagacgctagcgtga	o
683	gagaacgctggacctcatccat	o
684	gagaacgctagaccttctat	o
685	actagacgttagtgtga	o
686	cacaccttggtcaatgtcacgt	o
687	tctccatcctatggttttatcg	o
688	cgctggaccttccat	o
689	caccaccttggtcaatgtcacgt	o

- 52 -

690	gctagacgtagctgga	o
691	agtgcgattgcagatcg	o
692	ttttcgttttgtggtttgtggtt	
693	ttttcgttttgtcgttttgtcggtt	
694	ttttgtttttgtggtttgtggtt	
695	accgcatggattctaggcca	s
696	gctagacgtagcgt	o
697	aacgctggaccttccat	o
698	tcaazggt	o
699	ccttcgat	o
700	actagacgtagtgtga	s
701	gctagaggttagcgtga	s
702	atggactctccagcgttctc	o
703	atcgactctcgagcgttctc	o
704	gctagacgttagc	o
705	gctagacgt	o
706	agtgcgattcgagatcg	o
707	tcagzgct	o
708	ctgattgctctctcgta	o
709	tzaacggt	o
710	gagaaazgctggaccttccat	o
711	gctagacgttaggctga	o
712	gctacttagcgtga	o
713	gctaccttagcgtga	o
714	atcgacttcgagcgttctc	o
715	atgcactctgcagcgttctc	o
716	agtgactctccagcgttctc	o
717	gccagatgtagctgga	o
718	atcgactcgagcgttctc	o
719	atcgatcgagcgttctc	o
720	bgagaacgctcgaccttcgat	o
721	gctagacgtagctgga	sos
722	atcgactctcgagcgttctc	sos
723	tagacgttagcgtga	o
724	cgactctcgagcgttctc	o
725	ggggctcgaccttgagggggg	sos
726	gctaacgttagcgtga	o
727	cgtcgctgt	o
728	gagaacgctggacztccat	o
729	atcgacctacgtgcgttztc	o
730	atzgacctacgtgcgttctc	o
731	gctagazgttagcgt	o
732	atcgactctcgagzgttctc	o
733	ggggtaatgcatcagggggg	sos
734	ggctgtattcctgactgcc	s
735	ccatgctaacctctagc	o
736	gctagatgttagcgtga	o
737	cgtaccttacggtga	o
738	tccatgctggctcctgatgct	o
739	atcgactctctcgagcgttctc	o
740	gctagagcttagcgtga	o

- 53 -

741	atcgactctcgagtgttctc	o
742	aacgctcgaccttcgat	o
743	ctcaacgctggaccttccat	o
744	atcgacctacgtgcgttctc	o
745	gagaatgctggaccttccat	o
746	tcacgctaacctctgac	o
747	bgagaacgctccagcactgat	o
748	bgagcaagctggaccttccat	o
749	cgctagagggttagcgta	o
750	gctagatgttaacgt	o
751	atggaagggtccacgttctc	o
752	gctagatgttagcgt	o
753	gctagacgttagtgt	o
754	tccatgacgggtcctgatgct	o
755	tccatggcgggtcctgatgct	o
756	gctagacgatagcgt	o
757	gctagtcgatagcgt	o
758	tccatgacgttcctgatgct	o
759	tccatgtcggttcctgatgct	o
760	gctagacgttagzgt	o
761	gctaggcgttagcgt	o
762	tccatgtzgggtcctgatgct	o
763	tccatgtcggtzctgatgct	o
764	atzgactctzgagzgttctc	o
765	atggaagggtccagtgttctc	o
766	gcatgacgttgagct	o
767	gggggtcaacgttgagggggg	s
768	gggggtcaagtctgagggggg	sos
769	cgcgcgcgcgcgcgcgcgcg	o
770	cccccccccccccccccccccccc	s
771	cccccccccccccccccccccccc	s
772	tccatgtcgctcctgatcct	o
773	gctaaacgttagcgt	o
774	tccatgtcgatcctgatgct	o
775	tccatgccgggtcctgatgct	o
776	aaaatcaacgttgaaaaaaa	sos
777	tccataacgttcctgatgct	o
778	tggagggtcccaccgagatcggag	o
779	cgtcgtcgtcgtcgtcgtcg	s
780	ctgctgctgctgctgctgctg	s
781	gagaacgctccgaccttcgat	s
782	gctagatgttagcgt	s
783	gcatgacgttgagct	s
784	tcaatgctgaf	o
785	tcaacgttgaf	o
786	tcaacgttgab	o
787	gcaatattgcb	o
788	gcaatattgcf	o
789	agttgcaact	o
790	tcttcgaa	o
791	tcaacgtc	o

- 54 -

792	ccatgtcggtcctgatgct	o
793	gtttttatataatttggg	o
794	tttttggttgctgttttgctgt	o
795	ttgggggggggtt	s
796	gggggtgggggtt	s
797	ggtggtgtaggttttgg	o
798	bgagaazgctcgaccttcgat	o
799	tcaacgttaacgttaacgtt	o
800	bgagcaagztggaccttccat	o
801	bgagaazgctccagcactgat	o
802	tcaazgttgax	o
803	gzaatattgcx	o
804	tgtgtcttttgctgttttgctgt	o
805	ctgctgttagcaatttaactgtg	o
806	tccatgacgttcctgatgct	s
807	tgcattgccgtgcatccgtacacagctct	s
808	tgcattgccgtacacagctct	s
809	tgcattcagctct	s
810	tgcgtct	s
811	cccccccccccccccccc	s
812	cccccccccc	s
813	ccccccc	s
814	tgcattcagctct	sos
815	tgcattgccgtacacagctct	o
816	gagcaagctggaccttccat	s
817	tcaacgttaacgttaacgttaacgttaacgtt	s
818	gagaacgctcgaccttcgat	s
819	gtccccatttcccagaggaggaaat	o
820	ctagcggctgacgtcataaagctag	o
821	ctagcttgatgacgtcagccgctag	o
822	cggctgacgtcatcaa	s
823	ctgacgtg	o
824	ctgacgtcat	o
825	attcgatcggggcggggcgag	o
826	ctcgccccgccccgatcgaat	o
827	gactgacgtcagcgt	o
828	ctagcggctgacgtcataaagctagc	s
829	ctagctttatgacgtcagccgctagc	s
830	ctagcggctgagctcataaagctagc	s
831	ctagtggctgacgtcatcaagctag	s
832	tccaccacgtgggtctatgct	s
833	gggaatgaaagattttattataag	o
834	tctaaaaaccatctattcttaaccct	o
835	agctcaacgtcatgc	o
836	ttaacggtggtagcggatttggtc	o
837	ttaagaccaataccgctaccaccg	o
838	gatctagtgatgagtcagccggatc	o
839	gatccggctgactcatcactagatc	o
840	tccaagacgttcctgatgct	o
841	tccatgacgtccctgatgct	o
842	tccaccacgtgggtgatgct	o

- 55 -

843	ccacgtggacctctagc	o
844	tcagaccacgtggtcggtgttcctga	o
845	tcaggaacacccgaccacgtggtctga	o
846	catttccacgatttccca	o
847	ttcctctctgcaagagact	o
848	tgtatctctctgaaggact	o
849	ataaagcgaaactagcagcagtttc	o
850	gaaactgctgctagtttcgctttat	o
851	tgcccaaagaggaaaatttgtttcatacag	o
852	ctgtatgaaacaaattttcctctttgggca	o
853	ttaggggttaggggttaggggt	ss
854	tccatgagcttcctgatgct	ss
855	aaaacatgacgttcaaaaaa	ss
856	aaaacatgacgttcgggggg	ss
857	ggggcatgagcttcgggggg	sos
858	ctaggctgacgtcatcaagctagt	o
859	tctgacgtcatctgacgttggtgacgtct	o
860	ggaattagtaataagatatagaagt	o
861	ttacctttttataaacataactaaaacaaa	o
862	gcgtttttttttgcg	s
863	atatctaatacaaacattaacaaa	o
864	tctatcccaggtggttcctgtag	o
865	btccatgacgttcctgatgct	o
866	btccatgagcttcctgatgct	o
867	tttttttttttttf	o
868	tttttttttttttf	so
869	ctagcttgatgagctcagccgctag	o
870	ttcagttgtcttgctgcttagctaa	o
871	tccatgagcttcctgagct	s
872	ctagcggctgacgtcatcaatctag	o
873	tgctagctgtgcctgtacct	s
874	atgctaaaggacgtcacattgca	o
875	tgcaatgtgacgtccttttagcat	o
876	gtaggggactttccgagctcgagatcctatg	o
877	cataggatctcgagctcggaagtccttac	o
878	ctgtcaggaactgcaggtaagg	o
879	cataacataggaatatttactcctcgc	o
880	ctccagctccaagaaaggacg	o
881	gaagtttctggttaagtcttcg	o
882	tgctgcttttgtgcttttgtgctt	s
883	tcgtcggttttgtggttttgtggtt	s
884	tcgtcggttttgcgttttgcgtt	s
885	tcctgacgttcggcgcgccccc	s
886	tgctgcttttgtgcttttgtgctt	
887	tccatgagcttcctgagctt	s
888	tcgtcggtttcgctggttttgacgtt	s
889	tcgtcggtttgcgtgcgttttcgctgtt	s
890	tcgcgtgcgttttgcgttttgacgtt	s
891	ttcgctcggttttgcgttttgcgtt	s
892	tcctgacggggaagt	s
893	tcctggcgtggaagt	s

- 56 -

894	tcctggcgggtgaagt	s
895	tcctggcgttgaagt	s
896	tcctgacgtggaagt	s
897	gcgacgttcggcgcgcccc	s
898	gcgacgggcgcgcgcccc	s
899	gcggcgtcgggcgcgcccc	s
900	gcggcgtcgggcgcgcccc	s
901	gcgacgttcggcgcgcccc	s
902	gcggcgttcggcgcgcccc	s
903	gcgacgtcgggcgcgcccc	s
904	tcgtcgctgtctccg	s
905	tgtgggggttttggtttgg	s
906	aggggaggggaggggagggg	s
907	tgtgtgtgtgtgtgtgtgt	s
908	ctctctctctctctctctct	chimeric
909	ggggtcgacgtcgagggggg	s
910	atatatatatatatatatat	s
911	tttttttttttttttttttt	s
912	tttttttttttttttttttt	s
913	tttttttttttttttttttt	s
914	gctagaggggaggggt	
915	gctagatgttagggg	
916	gcatgagggggagct	
917	atggaagggtccagggggctc	
918	atggactctggagggggctc	
919	atggaagggtccaagggggctc	
920	gagaaggggggaccttgat	
921	gagaaggggggaccttccat	
922	gagaagggggccagcactgat	
923	tccatgtggggcctgatgct	
924	tccatgaggggcctgatgct	
925	tccatgtggggcctgctgat	
926	atggactctccggggttctc	
927	atggaagggtccggggttctc	
928	atggactctggaggggtctc	
929	atggagggtccatggggctc	
930	atggactctggggggttctc	
931	tccatgtgggtggggatgct	
932	tccatgcgggtggggatgct	
933	tccatgggggtcctgatgct	
934	tccatgggggtccctgatgct	
935	tccatgggggtgcctgatgct	
936	tccatgggggttccctgatgct	
937	tccatcgggggcctgatgct	
938	gctagagggagtggt	
939	tttttttttttttttttttt	s
940	gmgggtcaacgttgagggmggg	s
941	ggggagttcggttgaggggggg	s
942	tcgtcgtttccccccccccc	s
943	ttgggggggttttttttttttt	s
944	tttaaattttaaaatttaaata	s

- 57 -

945	ttgggttttttgggttttttttgg	s
946	tttcccttttcccttttccctc	s
947	ggggtcacgatgagggggg s	sos
948	tccatgacgttcctgacgtt	
949	tccatgacgttcctgacgtt	
950	tccatgacgttcctgacgtt	
951	tccatgacgttcctgacgtt	
952	tccatgacgttcctgacgtt	
953	tccatgacgttcctgacgtt	
954	tccatgacgttcctgacgtt	
955	tccatgacgttcctgacgtt	
956	tccatgacgttcctgacgtt	
957	tccatgacgttcctgacgtt	
958	tccatgacgttcctgacgtt	
959	gggggacgatcgtcggggg	sos
960	gggggtcgtacgacggggg	sos
961	ttttttttttttttttttttt	po
962	aaaaaaaaaaaaaaaaaaaaa	po
963	cccccccccccccccccccc	po
964	tcgtcgttttgcgttttgcgtt	
965	tcgtcgttttgcgttttgcgtt	
966	tcgtcgttttgcgttttgcgtt	
967	tcgtcgttttgcgttttgcgtt	
968	gggggtcaacgttgagggggg	
969	gggggtcaacgttgagggggg	
970	gggggtcaacgttgagggggg	
971	tgtgcttcccccccccccc	
972	ggggacgtcgacgtgggggg	sos
973	ggggtcgtcgacgagggggg	sos
974	ggggtcgacgtacgtcgaggggg	sos
975	ggggaccggtaccggtgggggg	sos
976	gggtcgacgtcgagggggg	sos
977	ggggtcgacgtcgagggggg	sos
978	ggggaacgttaacgttgggggg	sos
979	ggggtcaccggtgagggggg	sos
980	ggggtcgttcgaacgagggggg	sos
981	ggggacgttcgaacgtgggggg	sos
982	tcaactttga	s
983	tcaagcttga	s
984	tcacgatcgtga	s
985	tcagcatgctga	s
986	gggggagcatgctggggggg	sos
987	gggggggggggggggggggg	sos
988	gggggacgatatcgtcgggggg	sos
989	gggggacgacgtcgtcgggggg	sos
990	gggggacgagctcgtcgggggg	sos
991	gggggacgtacgtcgggggg	sos
992	tcaacgtt	
993	tccataccggtcctgatgct	
994	tccataccggtcctaccggt	s

- 58 -

995	gggggacgatcggtgggggg	sos
996	ggggaacgatcgtcgggggg	sos
997	ggg ggg acg atc gtc ggg ggg	sos
998	ggg gga cga tcg tcg ggg ggg	sos
999	aaa gac gtt aaa	po
1000	aaagagcttaaa	po
1001	aaagazgttaaa	po
1002	aaattcggaaaa	po
1003	gggggtcatcgatgagggggg	sos
1004	gggggtcaacggtgagggggg	sos
1005	atgtagcttaataacaaagc	po
1006	ggatcccttgagttacttct	po
1007	ccattccacttctgattacc	po
1008	tatgtattatcatgtagata	po
1009	agcctacgtattcaccctcc	po
1010	ttcctgcaactactattgta	po
1011	atagaaggccctacaccagt	po
1012	ttacaccggtctatggaggt	po
1013	ctaaccagatcaagtctagg	po
1014	cctagacttgatctggttag	po
1015	tataagcctcgccgacatg	po
1016	catgtcggacgaggttata	po
1017	tgggtggtgggagtaagctc	po
1018	gagctactccccaccacca	po
1019	gccttcgatcttcgttggga	po
1020	tggacttctctttgccgtct	po
1021	atgctgtagcccagcgataa	po
1022	accgaatcagcggaaagtga	po
1023	tccatgacgttctcgacgtt	
1024	ggagaaacccatgagctcatctgg	
1025	accacagaccagcaggcaga	
1026	gagcgtgaactgcgcgaaga	
1027	tcggtacccttgacgcggtt	
1028	ctggagccctagccaaggat	
1029	gcgactccatcaccagcgat	
1030	cctgaagtaagaaccagatgt	
1031	ctgtgttatctgacatacacc	
1032	aattagccttaggtgattggg	
1033	acatctggttcttacttcagg	
1034	ataagtcataatgttgggaactac	
1035	cccaatcacctaaggctaatt	
1036	ggggtcgtcgacgagggggg	sos
1037	ggggtcgttcgaacgagggggg	sos
1038	ggggacgttcgaacgtgggggg	sos
1039	tcctggcgqggaagt	s
1040	ggggaacgacgtcgttgggggg	sos
1041	ggggaacgtacgtcggggggg	sos
1042	ggggaacgtacgtacgttgggggg	sos
1043	ggggtcaccggtgagggggg	sos





- 60 -

1093	gggggttggttggttggtggggg	Sos
1094	tttttttttttttttttttttttttttttttt	
1095	aaaaaaaaaaaaaaaaaaaaaaaaaaaa	
1096	cccccccccccccccccccccccccccc	
1097	cgcgcgcgcgcgcgcgcgcgcgcgcg	

While CpG effects in mice are well characterized, information regarding the human system is limited. CpG phosphorothioate oligonucleotides with strong stimulatory activity in the mouse system show lower activity on human and other non-rodent immune cells. In the examples the development of a potent human CpG motif and the characterization of its effects and mechanisms of action on human primary B-cells is described. DNA containing this CpG motif strongly stimulated primary human B-cells to proliferate, to produce IL-6 and to express increased levels of CD86, CD40, CD54 and MHC II. It increased DNA binding activity of the transcription factors NFκB and AP-1, as well as phosphorylation of the stress activated protein kinases JNK and p38, and the transcription factor ATF-2. B-cell signaling pathways activated by CpG DNA were different from those activated by the B-cell receptor which activated ERK and a different isoform of JNK, but did not activate p38 and ATF-2. In general the data on CpG DNA-initiated signal transduction are consistent with those obtained in mice (Hacker H., et al. 1998. *Embo J* 17:6230, Yi A. K., and Krieg A. M. 1998. *J Immunol* 161:4493).

The preferred non-rodent motif is 5' TCGTCGTT 3'. Base exchanges within the most potent 8mer CpG motif (5' TCGTCGTT 3') diminished the activity of the oligonucleotide. The thymidines at the 5' and the 3' position of this motif were more important than the thymidine at the middle position. An adenine or guanosine at the middle position produced a decrease in the activity.

Of note, our studies demonstrate that one human CpG motif within a phosphodiester oligonucleotide (2080) is sufficient to produce the maximal effect, and that additional CpG motifs (2059) did not further enhance the activity. The oligonucleotide with the 8mer motif 5' TCG TCG TT 3' (2080) containing two CpG dinucleotides showed the highest activity in the studies. Replacement of the bases flanking the two CpG dinucleotides (5' position, middle position, 3' position) reduced the activity of this sequence. Both CpG dinucleotides within the 8mer CpG motif were required for the optimal activity (2108, 2106). Cytidine methylation of the CpG dinucleotides (2095) abolished the activity of 2080, while methylation of an unrelated

cytidine (2094) did not. The addition of two CpG motifs into the sequence of 2080 resulting in 2059 did not further increase the activity of the phosphodiester oligonucleotide. The sequence of 2080 with a phosphorothioate backbone (2116) demonstrated less activity, suggesting that additional CpG motifs are preferred for a  
5 potent phosphorothioate oligonucleotide.

It has been discovered according to the invention that the immunostimulatory nucleic acids have dramatic immune stimulatory effects on human cells such as NK cells, B cells, and DCs *in vitro*. It has been demonstrated that that the *in vitro* assays used herein predict *in vivo* effectiveness as a vaccine adjuvant in non-rodent vertebrates  
10 (Example 12), suggesting that immunostimulatory nucleic acids are effective therapeutic agents for human vaccination, cancer immunotherapy, asthma immunotherapy, general enhancement of immune function, enhancement of hematopoietic recovery following radiation or chemotherapy, and other immune modulatory applications.

Thus the immunostimulatory nucleic acids are useful in some aspects of the  
15 invention as a prophylactic vaccine for the treatment of a subject at risk of developing an infection with an infectious organism or a cancer in which a specific cancer antigen has been identified or an allergy or asthma where the allergen or predisposition to asthma is known. The immunostimulatory nucleic acids can also be given without the antigen or allergen for shorter term protection against infection, allergy or cancer, and in this case  
20 repeated doses will allow longer term protection. A subject at risk as used herein is a subject who has any risk of exposure to an infection causing pathogen or a cancer or an allergen or a risk of developing cancer. For instance, a subject at risk may be a subject who is planning to travel to an area where a particular type of infectious agent is found or it may be a subject who through lifestyle or medical procedures is exposed to bodily  
25 fluids which may contain infectious organisms or directly to the organism or even any subject living in an area where an infectious organism or an allergen has been identified. Subjects at risk of developing infection also include general populations to which a medical agency recommends vaccination with a particular infectious organism antigen. If the antigen is an allergen and the subject develops allergic responses to that particular  
30 antigen and the subject may be exposed to the antigen, i.e., during pollen season, then that subject is at risk of exposure to the antigen. A subject at risk of developing an allergy to asthma includes those subjects that have been identified as having an allergy or

- 62 -

asthma but that don't have the active disease during the immunostimulatory nucleic acid treatment as well as subjects that are considered to be at risk of developing these diseases because of genetic or environmental factors.

5 A subject at risk of developing a cancer is one who is who has a high probability of developing cancer. These subjects include, for instance, subjects having a genetic abnormality, the presence of which has been demonstrated to have a correlative relation to a higher likelihood of developing a cancer and subjects exposed to cancer causing agents such as tobacco, asbestos, or other chemical toxins, or a subject who has previously been treated for cancer and is in apparent remission. When a subject at risk of  
10 developing a cancer is treated with an antigen specific for the type of cancer to which the subject is at risk of developing and a immunostimulatory nucleic acid, the subject may be able to kill the cancer cells as they develop. If a tumor begins to form in the subject, the subject will develop a specific immune response against the tumor antigen.

In addition to the use of the immunostimulatory nucleic acids for prophylactic  
15 treatment, the invention also encompasses the use of the immunostimulatory nucleic acids for the treatment of a subject having an infection, an allergy, asthma, or a cancer.

A subject having an infection is a subject that has been exposed to an infectious pathogen and has acute or chronic detectable levels of the pathogen in the body. The immunostimulatory nucleic acids can be used with an antigen to mount an antigen  
20 specific systemic or mucosal immune response that is capable of reducing the level of or eradicating the infectious pathogen. An infectious disease, as used herein, is a disease arising from the presence of a foreign microorganism in the body. It is particularly important to develop effective vaccine strategies and treatments to protect the body's mucosal surfaces, which are the primary site of pathogenic entry.

25 A subject having an allergy is a subject that has or is at risk of developing an allergic reaction in response to an allergen. An allergy refers to acquired hypersensitivity to a substance (allergen). Allergic conditions include but are not limited to eczema, allergic rhinitis or coryza, hay fever, conjunctivitis, bronchial asthma, urticaria (hives) and food allergies, and other atopic conditions.

30 Currently, allergic diseases are generally treated by the injection of small doses of antigen followed by subsequent increasing dosage of antigen. It is believed that this procedure induces tolerization to the allergen to prevent further allergic reactions. These

- 63 -

methods, however, can take several years to be effective and are associated with the risk of side effects such as anaphylactic shock. The methods of the invention avoid these problems.

Allergies are generally caused by IgE antibody generation against harmless  
5 allergens. The cytokines that are induced by systemic or mucosal administration of immunostimulatory nucleic acids are predominantly of a class called Th1 (examples are IL-12 and IFN- $\gamma$ ) and these induce both humoral and cellular immune responses. The types of antibodies associated with a Th1 response are generally more protective because they have high neutralization and opsonization capabilities. The other major type of  
10 immune response, which is associated with the production of IL-4, IL-5 and IL-10 cytokines, is termed a Th2 immune response. Th2 responses involve predominately antibodies and these have less protective effect against infection and some Th2 isotypes (e.g., IgE) are associated with allergy. In general, it appears that allergic diseases are mediated by Th2 type immune responses while Th1 responses provide the best  
15 protection against infection, although excessive Th1 responses are associated with autoimmune disease. Based on the ability of the immunostimulatory nucleic acids to shift the immune response in a subject from a Th2 (which is associated with production of IgE antibodies and allergy) to a Th1 response (which is protective against allergic reactions), an effective dose for inducing an immune response of a immunostimulatory  
20 nucleic acid can be administered to a subject to treat or prevent an allergy.

Thus, the immunostimulatory nucleic acids have significant therapeutic utility in the treatment of allergic and non-allergic conditions such as asthma. Th2 cytokines, especially IL-4 and IL-5 are elevated in the airways of asthmatic subjects. These cytokines promote important aspects of the asthmatic inflammatory response, including  
25 IgE isotope switching, eosinophil chemotaxis and activation and mast cell growth. Th1 cytokines, especially IFN- $\gamma$  and IL-12, can suppress the formation of Th2 clones and production of Th2 cytokines. Asthma refers to a disorder of the respiratory system characterized by inflammation, narrowing of the airways and increased reactivity of the airways to inhaled agents. Asthma is frequently, although not exclusively associated  
30 with atopic or allergic symptoms.

A subject having a cancer is a subject that has detectable cancerous cells. The cancer may be a malignant or non-malignant cancer. Cancers or tumors include but are

- 64 -

not limited to biliary tract cancer; brain cancer; breast cancer; cervical cancer; choriocarcinoma; colon cancer; endometrial cancer; esophageal cancer; gastric cancer; intraepithelial neoplasms; lymphomas; liver cancer; lung cancer (e.g. small cell and non-small cell); melanoma; neuroblastomas; oral cancer; ovarian cancer; pancreas  
5 cancer; prostate cancer; rectal cancer; sarcomas; skin cancer; testicular cancer; thyroid cancer; and renal cancer, as well as other carcinomas and sarcomas. In one embodiment the cancer is hairy cell leukemia, chronic myelogenous leukemia, cutaneous T-cell leukemia, multiple myeloma, follicular lymphoma, malignant melanoma, squamous cell carcinoma, renal cell carcinoma, prostate carcinoma, bladder cell carcinoma, or colon  
10 carcinoma.

A subject according to the invention is a non-rodent subject. A non-rodent subject shall mean a human or vertebrate animal including but not limited to a dog, cat, horse, cow, pig, sheep, goat, chicken, primate, e.g., monkey, and fish (aquaculture species), e.g. salmon, but specifically excluding rodents such as rats and mice.

15 Thus, the invention can also be used to treat cancer and tumors in non human subjects. Cancer is one of the leading causes of death in companion animals (i.e., cats and dogs). Cancer usually strikes older animals which, in the case of house pets, have become integrated into the family. Forty-five % of dogs older than 10 years of age, are likely to succumb to the disease. The most common treatment options include surgery,  
20 chemotherapy and radiation therapy. Others treatment modalities which have been used with some success are laser therapy, cryotherapy, hyperthermia and immunotherapy. The choice of treatment depends on type of cancer and degree of dissemination. Unless the malignant growth is confined to a discrete area in the body, it is difficult to remove only malignant tissue without also affecting normal cells.

25 Malignant disorders commonly diagnosed in dogs and cats include but are not limited to lymphosarcoma, osteosarcoma, mammary tumors, mastocytoma, brain tumor, melanoma, adenosquamous carcinoma, carcinoid lung tumor, bronchial gland tumor, bronchiolar adenocarcinoma, fibroma, myxochondroma, pulmonary sarcoma, neurosarcoma, osteoma, papilloma, retinoblastoma, Ewing's sarcoma, Wilm's tumor,  
30 Burkitt's lymphoma, microglioma, neuroblastoma, osteoclastoma, oral neoplasia, fibrosarcoma, osteosarcoma and rhabdomyosarcoma. Other neoplasias in dogs include genital squamous cell carcinoma, transmissible venereal tumor, testicular tumor,

- 65 -

seminoma, Sertoli cell tumor, hemangiopericytoma, histiocytoma, chloroma (granulocytic sarcoma), corneal papilloma, corneal squamous cell carcinoma, hemangiosarcoma, pleural mesothelioma, basal cell tumor, thymoma, stomach tumor, adrenal gland carcinoma, oral papillomatosis, hemangioendothelioma and cystadenoma.

5 Additional malignancies diagnosed in cats include follicular lymphoma, intestinal lymphosarcoma, fibrosarcoma and pulmonary squamous cell carcinoma. The ferret, an ever-more popular house pet is known to develop insulinoma, lymphoma, sarcoma, neuroma, pancreatic islet cell tumor, gastric MALT lymphoma and gastric adenocarcinoma.

10 Neoplasias affecting agricultural livestock include leukemia, hemangiopericytoma and bovine ocular neoplasia (in cattle); preputial fibrosarcoma, ulcerative squamous cell carcinoma, preputial carcinoma, connective tissue neoplasia and mastocytoma (in horses); hepatocellular carcinoma (in swine); lymphoma and pulmonary adenomatosis (in sheep); pulmonary sarcoma, lymphoma, Rous sarcoma, 15 reticulendotheliosis, fibrosarcoma, nephroblastoma, B-cell lymphoma and lymphoid leukosis (in avian species); retinoblastoma, hepatic neoplasia, lymphosarcoma (lymphoblastic lymphoma), plasmacytoid leukemia and swimbladder sarcoma (in fish), caseous lymphadenitis (CLA): chronic, infectious, contagious disease of sheep and goats caused by the bacterium *Corynebacterium pseudotuberculosis*, and contagious lung 20 tumor of sheep caused by jaagsiekte.

The subject is exposed to the antigen. As used herein, the term exposed to refers to either the active step of contacting the subject with an antigen or the passive exposure of the subject to the antigen *in vivo*. Methods for the active exposure of a subject to an antigen are well-known in the art. In general, an antigen is administered directly to the 25 subject by any means such as intravenous, intramuscular, oral, transdermal, mucosal, intranasal, intratracheal, or subcutaneous administration. The antigen can be administered systemically or locally. Methods for administering the antigen and the immunostimulatory nucleic acid are described in more detail below. A subject is passively exposed to an antigen if an antigen becomes available for exposure to the 30 immune cells in the body. A subject may be passively exposed to an antigen, for instance, by entry of a foreign pathogen into the body or by the development of a tumor cell expressing a foreign antigen on its surface.

The methods in which a subject is passively exposed to an antigen can be particularly dependent on timing of administration of the immunostimulatory nucleic acid. For instance, in a subject at risk of developing a cancer or an infectious disease or an allergic or asthmatic response, the subject may be administered the  
5 immunostimulatory nucleic acid on a regular basis when that risk is greatest, i.e., during allergy season or after exposure to a cancer causing agent. Additionally the immunostimulatory nucleic acid may be administered to travelers before they travel to foreign lands where they are at risk of exposure to infectious agents. Likewise the immunostimulatory nucleic acid may be administered to soldiers or civilians at risk of  
10 exposure to biowarfare to induce a systemic or mucosal immune response to the antigen when and if the subject is exposed to it.

An antigen as used herein is a molecule capable of provoking an immune response. Antigens include but are not limited to cells, cell extracts, proteins, polypeptides, peptides, polysaccharides, polysaccharide conjugates, peptide and non-  
15 peptide mimics of polysaccharides and other molecules, small molecules, lipids, glycolipids, carbohydrates, viruses and viral extracts and multicellular organisms such as parasites and allergens. The term antigen broadly includes any type of molecule which is recognized by a host immune system as being foreign. Antigens include but are not limited to cancer antigens, microbial antigens, and allergens.

20 A cancer antigen as used herein is a compound, such as a peptide or protein, associated with a tumor or cancer cell surface and which is capable of provoking an immune response when expressed on the surface of an antigen presenting cell in the context of an MHC molecule. Cancer antigens can be prepared from cancer cells either by preparing crude extracts of cancer cells, for example, as described in Cohen, et al.,  
25 1994, *Cancer Research*, 54:1055, by partially purifying the antigens, by recombinant technology, or by de novo synthesis of known antigens. Cancer antigens include but are not limited to antigens that are recombinantly expressed, an immunogenic portion of, or a whole tumor or cancer. Such antigens can be isolated or prepared recombinantly or by any other means known in the art.

30 A microbial antigen as used herein is an antigen of a microorganism and includes but is not limited to virus, bacteria, parasites, and fungi. Such antigens include the intact microorganism as well as natural isolates and fragments or derivatives thereof and also



synthetic compounds which are identical to or similar to natural microorganism antigens and induce an immune response specific for that microorganism. A compound is similar to a natural microorganism antigen if it induces an immune response (humoral and/or cellular) to a natural microorganism antigen. Such antigens are used routinely in the art and are well known to those of ordinary skill in the art.

Examples of viruses that have been found in humans include but are not limited to: *Retroviridae* (e.g. human immunodeficiency viruses, such as HIV-1 (also referred to as HTLV-III, LAV or HTLV-III/LAV, or HIV-III; and other isolates, such as HIV-LP; *Picornaviridae* (e.g. polio viruses, hepatitis A virus; enteroviruses, human Coxsackie viruses, rhinoviruses, echoviruses); *Calciviridae* (e.g. strains that cause gastroenteritis); *Togaviridae* (e.g. equine encephalitis viruses, rubella viruses); *Flaviridae* (e.g. dengue viruses, encephalitis viruses, yellow fever viruses); *Coronaviridae* (e.g. coronaviruses); *Rhabdoviridae* (e.g. vesicular stomatitis viruses, rabies viruses); *Coronaviridae* (e.g. coronaviruses); *Rhabdoviridae* (e.g. vesicular stomatitis viruses, rabies viruses); *Filoviridae* (e.g. ebola viruses); *Paramyxoviridae* (e.g. parainfluenza viruses, mumps virus, measles virus, respiratory syncytial virus); *Orthomyxoviridae* (e.g. influenza viruses); *Bunyaviridae* (e.g. Hantaan viruses, bunya viruses, phleboviruses and Nairo viruses); *Arenaviridae* (hemorrhagic fever viruses); *Reoviridae* (e.g. reoviruses, orbiviruses and rotaviruses); *Birnaviridae*; *Hepadnaviridae* (Hepatitis B virus); *Parvoviridae* (parvoviruses); *Papovaviridae* (papilloma viruses, polyoma viruses); *Adenoviridae* (most adenoviruses); *Herpesviridae* (herpes simplex virus (HSV) 1 and 2, varicella zoster virus, cytomegalovirus (CMV), herpes virus; *Poxviridae* (variola viruses, vaccinia viruses, pox viruses); and *Iridoviridae* (e.g. African swine fever virus); and unclassified viruses (e.g. the etiological agents of Spongiform encephalopathies, the agent of delta hepatitis (thought to be a defective satellite of hepatitis B virus), the agents of non-A, non-B hepatitis (class 1 = internally transmitted; class 2 = parenterally transmitted (i.e. Hepatitis C); Norwalk and related viruses, and astroviruses).

Both gram negative and gram positive bacteria serve as antigens in vertebrate animals. Such gram positive bacteria include, but are not limited to, *Pasteurella* species, *Staphylococci* species, and *Streptococcus* species. Gram negative bacteria include, but are not limited to, *Escherichia coli*, *Pseudomonas* species, and *Salmonella* species. Specific examples of infectious bacteria include but are not limited to, *Helicobacter*

*pyloris*, *Borelia burgdorferi*, *Legionella pneumophila*, *Mycobacteria* *sps* (e.g. *M. tuberculosis*, *M. avium*, *M. intracellulare*, *M. kansasii*, *M. gordonae*), *Staphylococcus aureus*, *Neisseria gonorrhoeae*, *Neisseria meningitidis*, *Listeria monocytogenes*, *Streptococcus pyogenes* (Group A Streptococcus), *Streptococcus agalactiae* (Group B Streptococcus), *Streptococcus* (viridans group), *Streptococcus faecalis*, *Streptococcus bovis*, *Streptococcus* (anaerobic *sps.*), *Streptococcus pneumoniae*, pathogenic *Campylobacter sp.*, *Enterococcus sp.*, *Haemophilus influenzae*, *Bacillus anthracis*, *Corynebacterium diphtheriae*, *Corynebacterium sp.*, *Erysipelothrix rhusiopathiae*, *Clostridium perfringens*, *Clostridium tetani*, *Enterobacter aerogenes*, *Klebsiella pneumoniae*, *Pasturella multocida*, *Bacteroides sp.*, *Fusobacterium nucleatum*, *Streptobacillus moniliformis*, *Treponema pallidum*, *Treponema pertenue*, *Leptospira*, *Rickettsia*, and *Actinomyces israeli*.

Examples of fungi include *Cryptococcus neoformans*, *Histoplasma capsulatum*, *Coccidioides immitis*, *Blastomyces dermatitidis*, *Chlamydia trachomatis*, *Candida albicans*.

Other infectious organisms (i.e., protists) include *Plasmodium spp.* such as *Plasmodium falciparum*, *Plasmodium malariae*, *Plasmodium ovale*, and *Plasmodium vivax* and *Toxoplasma gondii*. Blood-borne and/or tissues parasites include *Plasmodium spp.*, *Babesia microti*, *Babesia divergens*, *Leishmania tropica*, *Leishmania spp.*, *Leishmania braziliensis*, *Leishmania donovani*, *Trypanosoma gambiense* and *Trypanosoma rhodesiense* (African sleeping sickness), *Trypanosoma cruzi* (Chagas' disease), and *Toxoplasma gondii*.

Other medically relevant microorganisms have been described extensively in the literature, e.g., see C.G.A Thomas, *Medical Microbiology*, Bailliere Tindall, Great Britain 1983, the entire contents of which is hereby incorporated by reference.

Although many of the microbial antigens described above relate to human disorders, the invention is also useful for treating other nonhuman vertebrates. Nonhuman vertebrates are also capable of developing infections which can be prevented or treated with the Immunostimulatory nucleic acids disclosed herein. For instance, in addition to the treatment of infectious human diseases, the methods of the invention are useful for treating infections of animals.

- 69 -

As used herein, the term treat, treated, or treating when used with respect to an infectious disease refers to a prophylactic treatment which increases the resistance of a subject (a subject at risk of infection) to infection with a pathogen or, in other words, decreases the likelihood that the subject will become infected with the pathogen as well  
5 as a treatment after the subject (a subject who has been infected) has become infected in order to fight the infection, e.g., reduce or eliminate the infection or prevent it from becoming worse.

Many vaccines for the treatment of non-human vertebrates are disclosed in Bennett, K. *Compendium of Veterinary Products*, 3rd ed. North American  
10 Compendiums, Inc., 1995. As discussed above, antigens include infectious microbes such as virus, parasite, bacteria and fungi and fragments thereof, derived from natural sources or synthetically. Infectious viruses of both human and non-human vertebrates, include retroviruses, RNA viruses and DNA viruses. This group of retroviruses includes both simple retroviruses and complex retroviruses. The simple retroviruses include the  
15 subgroups of B-type retroviruses, C-type retroviruses and D-type retroviruses. An example of a B-type retrovirus is mouse mammary tumor virus (MMTV). The C-type retroviruses include subgroups C-type group A (including Rous sarcoma virus (RSV), avian leukemia virus (ALV), and avian myeloblastosis virus (AMV)) and C-type group B (including feline leukemia virus (FeLV), gibbon ape leukemia virus (GALV), spleen  
20 necrosis virus (SNV), reticuloendotheliosis virus (RV) and simian sarcoma virus (SSV)). The D-type retroviruses include Mason-Pfizer monkey virus (MPMV) and simian retrovirus type 1 (SRV-1). The complex retroviruses include the subgroups of lentiviruses, T-cell leukemia viruses and the foamy viruses. Lentiviruses include HIV-1, but also include HIV-2, SIV, Visna virus, feline immunodeficiency virus (FIV), and  
25 equine infectious anemia virus (EIAV). The T-cell leukemia viruses include HTLV-1, HTLV-II, simian T-cell leukemia virus (STLV), and bovine leukemia virus (BLV). The foamy viruses include human foamy virus (HFV), simian foamy virus (SFV) and bovine foamy virus (BFV).

Examples of other RNA viruses that are antigens in vertebrate animals include,  
30 but are not limited to, members of the family Reoviridae, including the genus Orthoreovirus (multiple serotypes of both mammalian and avian retroviruses), the genus Orbivirus (Bluetongue virus, Eugenangee virus, Kemerovo virus, African horse sickness

- 70 -

virus, and Colorado Tick Fever virus), the genus Rotavirus (human rotavirus, Nebraska calf diarrhea virus, simian rotavirus, bovine or ovine rotavirus, avian rotavirus); the family Picornaviridae, including the genus Enterovirus (poliovirus, Coxsackie virus A and B, enteric cytopathic human orphan (ECHO) viruses, hepatitis A virus, Simian  
5 enteroviruses, Murine encephalomyelitis (ME) viruses, Poliovirus muris, Bovine enteroviruses, Porcine enteroviruses, the genus Cardiovirus (Encephalomyocarditis virus (EMC), Mengovirus), the genus Rhinovirus (Human rhinoviruses including at least 113 subtypes; other rhinoviruses), the genus Aphthovirus (Foot and Mouth disease (FMDV); the family Calciviridae, including Vesicular exanthema of swine virus, San Miguel sea  
10 lion virus, Feline picornavirus and Norwalk virus; the family Togaviridae, including the genus Alphavirus (Eastern equine encephalitis virus, Semliki forest virus, Sindbis virus, Chikungunya virus, O'Nyong-Nyong virus, Ross river virus, Venezuelan equine encephalitis virus, Western equine encephalitis virus), the genus Flavivirus (Mosquito borne yellow fever virus, Dengue virus, Japanese encephalitis virus, St. Louis  
15 encephalitis virus, Murray Valley encephalitis virus, West Nile virus, Kunjin virus, Central European tick borne virus, Far Eastern tick borne virus, Kyasanur forest virus, Louping III virus, Powassan virus, Omsk hemorrhagic fever virus), the genus Rubivirus (Rubella virus), the genus Pestivirus (Mucosal disease virus, Hog cholera virus, Border disease virus); the family Bunyaviridae, including the genus Bunyavirus (Bunyamwera  
20 and related viruses, California encephalitis group viruses), the genus Phlebovirus (Sandfly fever Sicilian virus, Rift Valley fever virus), the genus Nairovirus (Crimean-Congo hemorrhagic fever virus, Nairobi sheep disease virus), and the genus Uukuvirus (Uukuniemi and related viruses); the family Orthomyxoviridae, including the genus Influenza virus (Influenza virus type A, many human subtypes); Swine influenza  
25 virus, and Avian and Equine Influenza viruses; influenza type B (many human subtypes), and influenza type C (possible separate genus); the family paramyxoviridae, including the genus Paramyxovirus (Parainfluenza virus type 1, Sendai virus, Hemadsorption virus, Parainfluenza viruses types 2 to 5, Newcastle Disease Virus, Mumps virus), the genus Morbillivirus (Measles virus, subacute sclerosing panencephalitis virus, distemper virus,  
30 Rinderpest virus), the genus Pneumovirus (respiratory syncytial virus (RSV), Bovine respiratory syncytial virus and Pneumonia virus); the family Rhabdoviridae, including the genus Vesiculovirus (VSV), Chandipura virus, Flanders-Hart Park virus), the genus

Lyssavirus (Rabies virus), fish Rhabdoviruses, and two probable Rhabdoviruses (Marburg virus and Ebola virus); the family Arenaviridae, including Lymphocytic choriomeningitis virus (LCM), Tacaribe virus complex, and Lassa virus; the family Coronaviridae, including Infectious Bronchitis Virus (IBV), Hepatitis virus, Human enteric corona virus, and Feline infectious peritonitis (Feline coronavirus).

Illustrative DNA viruses that are antigens in vertebrate animals include, but are not limited to, the family Poxviridae, including the genus Orthopoxvirus (Variola major, Variola minor, Monkey pox Vaccinia, Cowpox, Buffalopox, Rabbitpox, Ectromelia), the genus Leporipoxvirus (Myxoma, Fibroma), the genus Avipoxvirus (Fowlpox, other avian poxvirus), the genus Capripoxvirus (sheeppox, goatpox), the genus Suipoxvirus (Swinepox), the genus Parapoxvirus (contagious postular dermatitis virus, pseudocowpox, bovine papular stomatitis virus); the family Iridoviridae (African swine fever virus, Frog viruses 2 and 3, Lymphocystis virus of fish); the family Herpesviridae, including the alpha-Herpesviruses (Herpes Simplex Types 1 and 2, Varicella-Zoster, Equine abortion virus, Equine herpes virus 2 and 3, pseudorabies virus, infectious bovine keratoconjunctivitis virus, infectious bovine rhinotracheitis virus, feline rhinotracheitis virus, infectious laryngotracheitis virus) the Beta-herpesviruses (Human cytomegalovirus and cytomegaloviruses of swine and monkeys); the gamma-herpesviruses (Epstein-Barr virus (EBV), Marek's disease virus, Herpes saimiri, Herpesvirus ateles, Herpesvirus sylvilagus, guinea pig herpes virus, Lucke tumor virus); the family Adenoviridae, including the genus Mastadenovirus (Human subgroups A,B,C,D,E and ungrouped; simian adenoviruses (at least 23 serotypes), infectious canine hepatitis, and adenoviruses of cattle, pigs, sheep, frogs and many other species, the genus Aviadenovirus (Avian adenoviruses); and non-cultivable adenoviruses; the family Papoviridae, including the genus Papillomavirus (Human papilloma viruses, bovine papilloma viruses, Shope rabbit papilloma virus, and various pathogenic papilloma viruses of other species), the genus Polyomavirus (polyomavirus, Simian vacuolating agent (SV-40), Rabbit vacuolating agent (RKV), K virus, BK virus, JC virus, and other primate polyoma viruses such as Lymphotropic papilloma virus); the family Parvoviridae including the genus Adeno-associated viruses, the genus Parvovirus (Feline panleukopenia virus, bovine parvovirus, canine parvovirus, Aleutian mink disease virus, etc). Finally, DNA viruses may include viruses which do not fit into the above families such as Kuru and

- 72 -

Creutzfeldt-Jacob disease viruses and chronic infectious neuropathic agents (CHINA virus).

Each of the foregoing lists is illustrative, and is not intended to be limiting.

In addition to the use of the immunostimulatory nucleic acids to induce an  
5 antigen specific immune response in humans, the methods of the preferred embodiments are particularly well suited for treatment of birds such as hens, chickens, turkeys, ducks, geese, quail, and pheasant. Birds are prime targets for many types of infections.

Hatching birds are exposed to pathogenic microorganisms shortly after birth. Although these birds are initially protected against pathogens by maternal derived  
10 antibodies, this protection is only temporary, and the bird's own immature immune system must begin to protect the bird against the pathogens. It is often desirable to prevent infection in young birds when they are most susceptible. It is also desirable to prevent against infection in older birds, especially when the birds are housed in closed quarters, leading to the rapid spread of disease. Thus, it is desirable to administer the  
15 Immunostimulatory nucleic acid and the non-nucleic acid adjuvant of the invention to birds to enhance an antigen-specific immune response when antigen is present.

An example of a common infection in chickens is chicken infectious anemia virus (CIAV). CIAV was first isolated in Japan in 1979 during an investigation of a Marek's disease vaccination break (Yuasa et al., 1979, Avian Dis. 23:366-385). Since  
20 that time, CIAV has been detected in commercial poultry in all major poultry producing countries (van Bulow et al., 1991, pp.690-699) in Diseases of Poultry, 9th edition, Iowa State University Press).

CIAV infection results in a clinical disease, characterized by anemia, hemorrhage and immunosuppression, in young susceptible chickens. Atrophy of the  
25 thymus and of the bone marrow and consistent lesions of CIAV-infected chickens are also characteristic of CIAV infection. Lymphocyte depletion in the thymus, and occasionally in the bursa of Fabricius, results in immunosuppression and increased susceptibility to secondary viral, bacterial, or fungal infections which then complicate the course of the disease. The immunosuppression may cause aggravated disease after  
30 infection with one or more of Marek's disease virus (MDV), infectious bursal disease virus, reticuloendotheliosis virus, adenovirus, or reovirus. It has been reported that pathogenesis of MDV is enhanced by CIAV (DeBoer et al., 1989, p. 28 In Proceedings

- 73 -

of the 38th Western Poultry Diseases Conference, Tempe, Ariz.). Further, it has been reported that CIAV aggravates the signs of infectious bursal disease (Rosenberger et al., 1989, Avian Dis. 33:707-713). Chickens develop an age resistance to experimentally induced disease due to CAA. This is essentially complete by the age of 2 weeks, but  
5 older birds are still susceptible to infection (Yuasa, N. et al., 1979 supra; Yuasa, N. et al., Arian Diseases 24, 202-209, 1980). However, if chickens are dually infected with CAA and an immunosuppressive agent (IBDV, MDV etc.), age resistance against the disease is delayed (Yuasa, N. et al., 1979 and 1980 supra; Bulow von V. et al., J. Veterinary Medicine 33, 93-116, 1986). Characteristics of CIAV that may potentiate disease  
10 transmission include high resistance to environmental inactivation and some common disinfectants. The economic impact of CIAV infection on the poultry industry is clear from the fact that 10% to 30% of infected birds in disease outbreaks die.

Vaccination of birds, like other vertebrate animals can be performed at any age. Normally, vaccinations are performed at up to 12 weeks of age for a live microorganism  
15 and between 14-18 weeks for an inactivated microorganism or other type of vaccine. For in ovo vaccination, vaccination can be performed in the last quarter of embryo development. The vaccine may be administered subcutaneously, by spray, orally, intraocularly, intratracheally, nasally, or by other mucosal delivery methods described herein. Thus, the immunostimulatory nucleic acids of the invention can be administered  
20 to birds and other non-human vertebrates using routine vaccination schedules and the antigen can be administered after an appropriate time period as described herein.

Cattle and livestock are also susceptible to infection. Diseases which affect these animals can produce severe economic losses, especially amongst cattle. The methods of the invention can be used to protect against infection in livestock, such as cows, horses,  
25 pigs, sheep, and goats.

Cows can be infected by bovine viruses. Bovine viral diarrhea virus (BVDV) is a small enveloped positive-stranded RNA virus and is classified, along with hog cholera virus (HOCV) and sheep border disease virus (BDV), in the pestivirus genus. Although, Pestiviruses were previously classified in the Togaviridae family, some studies have  
30 suggested their reclassification within the Flaviviridae family along with the flavivirus and hepatitis C virus (HCV) groups (Francki, et al., 1991).

- 74 -

BVDV, which is an important pathogen of cattle can be distinguished, based on cell culture analysis, into cytopathogenic (CP) and noncytopathogenic (NCP) biotypes. The NCP biotype is more widespread although both biotypes can be found in cattle. If a pregnant cow becomes infected with an NCP strain, the cow can give birth to a persistently infected and specifically immunotolerant calf that will spread virus during its lifetime. The persistently infected cattle can succumb to mucosal disease and both biotypes can then be isolated from the animal. Clinical manifestations can include abortion, teratogenesis, and respiratory problems, mucosal disease and mild diarrhea. In addition, severe thrombocytopenia, associated with herd epidemics, that may result in the death of the animal has been described and strains associated with this disease seem more virulent than the classical BVDVs.

Equine herpes viruses (EHV) comprise a group of antigenically distinct biological agents which cause a variety of infections in horses ranging from subclinical to fatal disease. These include Equine herpesvirus-1 (EHV-1), a ubiquitous pathogen in horses. EHV-1 is associated with epidemics of abortion, respiratory tract disease, and central nervous system disorders. Primary infection of upper respiratory tract of young horses results in a febrile illness which lasts for 8 to 10 days. Immunologically experienced mares may be re-infected via the respiratory tract without disease becoming apparent, so that abortion usually occurs without warning. The neurological syndrome is associated with respiratory disease or abortion and can affect animals of either sex at any age, leading to lack of co-ordination, weakness and posterior paralysis (Telford, E. A. R. et al., Virology 189, 304-316, 1992). Other EHV's include EHV-2, or equine cytomegalovirus, EHV-3, equine coital exanthema virus, and EHV-4, previously classified as EHV-1 subtype 2.

Sheep and goats can be infected by a variety of dangerous microorganisms including visna-maedi.

Primates such as monkeys, apes and macaques can be infected by simian immunodeficiency virus. Inactivated cell-virus and cell-free whole simian immunodeficiency vaccines have been reported to afford protection in macaques (Stott et al. (1990) Lancet 36:1538-1541; Desrosiers et al. PNAS USA (1989) 86:6353-6357; Murphey-Corb et al. (1989) Science 246:1293-1297; and Carlson et al. (1990) AIDS Res. Human Retroviruses 6:1239-1246). A recombinant HIV gp120 vaccine has been



- 75 -

reported to afford protection in chimpanzees (Berman et al. (1990) Nature 345:622-625).

Cats, both domestic and wild, are susceptible to infection with a variety of microorganisms. For instance, feline infectious peritonitis is a disease which occurs in both domestic and wild cats, such as lions, leopards, cheetahs, and jaguars. When it is desirable to prevent infection with this and other types of pathogenic organisms in cats, the methods of the invention can be used to vaccinate cats to protect them against infection.

Domestic cats may become infected with several retroviruses, including but not limited to feline leukemia virus (FeLV), feline sarcoma virus (FeSV), endogenous type Concornavirus (RD-114), and feline syncytia-forming virus (FeSFV). Of these, FeLV is the most significant pathogen, causing diverse symptoms, including lymphoreticular and myeloid neoplasms, anemias, immune mediated disorders, and an immunodeficiency syndrome which is similar to human acquired immune deficiency syndrome (AIDS). Recently, a particular replication-defective FeLV mutant, designated FeLV-AIDS, has been more particularly associated with immunosuppressive properties.

The discovery of feline T-lymphotropic lentivirus (also referred to as feline immunodeficiency) was first reported in Pedersen et al. (1987) Science 235:790-793. Characteristics of FIV have been reported in Yamamoto et al. (1988) Leukemia, December Supplement 2:204S-215S; Yamamoto et al. (1988) Am. J. Vet. Res. 49:1246-1258; and Ackley et al. (1990) J. Virol. 64:5652-5655. Cloning and sequence analysis of FIV have been reported in Olmsted et al. (1989) Proc. Natl. Acad. Sci. USA 86:2448-2452 and 86:4355-4360.

Feline infectious peritonitis (FIP) is a sporadic disease occurring unpredictably in domestic and wild Felidae. While FIP is primarily a disease of domestic cats, it has been diagnosed in lions, mountain lions, leopards, cheetahs, and the jaguar. Smaller wild cats that have been afflicted with FIP include the lynx and caracal, sand cat, and pallas cat. In domestic cats, the disease occurs predominantly in young animals, although cats of all ages are susceptible. A peak incidence occurs between 6 and 12 months of age. A decline in incidence is noted from 5 to 13 years of age, followed by an increased incidence in cats 14 to 15 years old.

- 76 -

Viral, bacterial, and parasitic diseases in fin-fish, shellfish or other aquatic life forms pose a serious problem for the aquaculture industry. Owing to the high density of animals in the hatchery tanks or enclosed marine farming areas, infectious diseases may eradicate a large proportion of the stock in, for example, a fin-fish, shellfish, or other aquatic life forms facility. Prevention of disease is a more desired remedy to these threats to fish than intervention once the disease is in progress. Vaccination of fish is the only preventative method which may offer long-term protection through immunity. Nucleic acid based vaccinations are described in US Patent No. 5,780,448 issued to Davis.

The fish immune system has many features similar to the mammalian immune system, such as the presence of B cells, T cells, lymphokines, complement, and immunoglobulins. Fish have lymphocyte subclasses with roles that appear similar in many respects to those of the B and T cells of mammals. Vaccines can be administered by immersion or orally.

Aquaculture species include but are not limited to fin-fish, shellfish, and other aquatic animals. Fin-fish include all vertebrate fish, which may be bony or cartilaginous fish, such as, for example, salmonids, carp, catfish, yellowtail, seabream, and seabass. Salmonids are a family of fin-fish which include trout (including rainbow trout), salmon, and Arctic char. Examples of shellfish include, but are not limited to, clams, lobster, shrimp, crab, and oysters. Other cultured aquatic animals include, but are not limited to eels, squid, and octopi.

Polypeptides of viral aquaculture pathogens include but are not limited to glycoprotein (G) or nucleoprotein (N) of viral hemorrhagic septicemia virus (VHSV); G or N proteins of infectious hematopoietic necrosis virus (IHNV); VP1, VP2, VP3 or N structural proteins of infectious pancreatic necrosis virus (IPNV); G protein of spring viremia of carp (SVC); and a membrane-associated protein, tegumin or capsid protein or glycoprotein of channel catfish virus (CCV).

Typical parasites infecting horses are *Gasterophilus* spp.; *Eimeria leuckarti*, *Giardia* spp.; *Trichostrongylus axei*; *Babesia* spp. (RBC's), *Theileria equi*; *Trypanosoma* spp.; *Klossiella equi*; *Sarcocystis* spp.

- 77 -

Typical parasites infecting swine include *Eimeria bebliecki*, *Eimeria scabra*, *Isospora suis*, *Giardia* spp.; *Balantidium coli*, *Entamoeba histolytica*; *Toxoplasma gondii* and *Sarcocystis* spp., and *Trichinella spiralis*.

The major parasites of dairy and beef cattle include *Eimeria* spp.,  
 5 *Cryptosporidium* sp., *Giardia* spp.; *Toxoplasma gondii*; *Babesia bovis* (RBC), *Babesia bigemina* (RBC), *Trypanosoma* spp. (plasma), *Theileria* spp. (RBC); *Theileria parva* (lymphocytes); *Tritrichomonas foetus*; and *Sarcocystis* spp.

The major parasites of raptors include *Trichomonas gallinae*; Coccidia (*Eimeria* spp.); *Plasmodium relictum*, *Leucocytozoon danilewskyi* (owls), *Haemoproteus* spp.,  
 10 *Trypanosoma* spp.; *Histomonas*; *Cryptosporidium meleagridis*, *Cryptosporidium baileyi*, *Giardia*, *Eimeria*; *Toxoplasma*.

Typical parasites infecting sheep and goats include *Eimeria* spp., *Cryptosporidium* sp., *Giardia* sp.; *Toxoplasma gondii*; *Babesia* spp. (RBC), *Trypanosoma* spp. (plasma), *Theileria* spp. (RBC); and *Sarcocystis* spp.

15 Typical parasitic infections in poultry include coccidiosis caused by *Eimeria acervulina*, *E. necatrix*, *E. tenella*, *Isospora* spp. and *Eimeria truncata*; histomoniasis, caused by *Histomonas meleagridis* and *Histomonas gallinarum*; trichomoniasis caused by *Trichomonas gallinae*; and hexamitiasis caused by *Hexamita meleagridis*. Poultry can also be infected *Emeria maxima*, *Emeria meleagridis*, *Eimeria adenoeides*, *Eimeria*  
 20 *meleagrimitis*, *Cryptosporidium*, *Eimeria brunetti*, *Emeria adenoeides*, *Leucocytozoon* spp., *Plasmodium* spp., *Hemoproteus meleagridis*, *Toxoplasma gondii* and *Sarcocystis*.

The methods of the invention can also be applied to the treatment and/or prevention of parasitic infection in dogs, cats, birds, fish and ferrets. Typical parasites of birds include *Trichomonas gallinae*; *Eimeria* spp., *Isospora* spp., *Giardia*;  
 25 *Cryptosporidium*; *Sarcocystis* spp., *Toxoplasma gondii*, *Haemoproteus/Parahaemoproteus*, *Plasmodium* spp., *Leucocytozoon/Akiba*, *Atoxoplasma*, *Trypanosoma* spp. Typical parasites infecting dogs include *Trichinella spiralis*; *Isopora* spp., *Sarcocystis* spp., *Cryptosporidium* spp., *Hammondia* spp., *Giardia duodenalis (canis)*; *Balantidium coli*, *Entamoeba histolytica*; *Hepatozoon canis*;  
 30 *Toxoplasma gondii*, *Trypanosoma cruzi*; *Babesia canis*; *Leishmania amastigotes*; *Neospora caninum*.

Typical parasites infecting feline species include *Isospora* spp., *Toxoplasma gondii*, *Sarcocystis* spp., *Hammondia hammondi*, *Besnoitia* spp., *Giardia* spp.; *Entamoeba histolytica*; *Hepatozoon canis*, *Cytauxzoon* sp., *Cytauxzoon* sp., *Cytauxzoon* sp. (red cells, RE cells).

- 5        Typical parasites infecting fish include *Hexamita* spp., *Eimeria* spp.; *Cryptobia* spp., *Nosema* spp., *Myxosoma* spp., *Chilodonella* spp., *Trichodina* spp.; *Plistophora* spp., *Myxosoma Henneguya*, *Costia* spp., *Ichthyophthirius* spp., and *Oodinium* spp.

- Typical parasites of wild mammals include *Giardia* spp. (carnivores, herbivores), *Isospora* spp. (carnivores), *Eimeria* spp. (carnivores, herbivores); *Theileria* spp. (herbivores), *Babesia* spp. (carnivores, herbivores), *Trypanosoma* spp. (carnivores, herbivores); *Schistosoma* spp. (herbivores); *Fasciola hepatica* (herbivores), *Fascioloides magna* (herbivores), *Fasciola gigantica* (herbivores), *Trichinella spiralis* (carnivores, herbivores).
- 10

- Parasitic infections in zoos can also pose serious problems. Typical parasites of the bovidae family (blesbok, antelope, banteng, eland, gaur, impala, klipspringer, kudu, gazelle) include *Eimeria* spp. Typical parasites in the pinnipedae family (seal, sea lion) include *Eimeria phocae*. Typical parasites in the camelidae family (camels, llamas) include *Eimeria* spp. Typical parasites of the giraffidae family (giraffes) include *Eimeria* spp. Typical parasites in the elephantidae family (African and Asian) include *Fasciola* spp. Typical parasites of lower primates (chimpanzees, orangutans, apes, baboons, macaques, monkeys) include *Giardia* sp.; *Balantidium coli*, *Entamoeba histolytica*, *Sarcocystis* spp., *Toxoplasma gondii*; *Plasmodium* spp. (RBC), *Babesia* spp. (RBC), *Trypanosoma* spp. (plasma), *Leishmania* spp. (macrophages).
- 15
- 20

- Polypeptides of bacterial pathogens include but are not limited to an iron-regulated outer membrane protein, (IROMP), an outer membrane protein (OMP), and an A-protein of *Aeromonis salmonicida* which causes furunculosis, p57 protein of *Renibacterium salmoninarum* which causes bacterial kidney disease (BKD), major surface associated antigen (msa), a surface expressed cytotoxin (mpr), a surface expressed hemolysin (ish), and a flagellar antigen of *Yersinia*; an extracellular protein (ECP), an iron-regulated outer membrane protein (IROMP), and a structural protein of *Pasteurellosis*; an OMP and a flagellar protein of *Vibrio anguillarum* and *V. ordalii*; a flagellar protein, an OMP protein, *aroA*, and *purA* of *Edwardsiella ictaluri* and *E.*
- 25
- 30

tarda; and surface antigen of *Ichthyophthirius*; and a structural and regulatory protein of *Cytophaga columnari*; and a structural and regulatory protein of *Rickettsia*.

Polypeptides of a parasitic pathogen include but are not limited to the surface antigens of *Ichthyophthirius*.

- 5       An allergen refers to a substance (antigen) that can induce an allergic or asthmatic response in a susceptible subject. The list of allergens is enormous and can include pollens, insect venoms, animal dander dust, fungal spores and drugs (e.g. penicillin). Examples of natural, animal and plant allergens include but are not limited to proteins specific to the following genres: *Canine* (*Canis familiaris*); *Dermatophagoides* (e.g. *Dermatophagoides farinae*); *Felis* (*Felis domesticus*); *Ambrosia* (*Ambrosia artemisiifolia*); *Lolium* (e.g. *Lolium perenne* or *Lolium multiflorum*); *Cryptomeria* (*Cryptomeria japonica*); *Alternaria* (*Alternaria alternata*); *Alder*; *Alnus* (*Alnus gultinoasa*); *Betula* (*Betula verrucosa*); *Quercus* (*Quercus alba*); *Olea* (*Olea europa*); *Artemisia* (*Artemisia vulgaris*); *Plantago* (e.g. *Plantago lanceolata*); *Parietaria* (e.g. *Parietaria officinalis* or *Parietaria judaica*); *Blattella* (e.g. *Blattella germanica*); *Apis* (e.g. *Apis multiflorum*); *Cupressus* (e.g. *Cupressus sempervirens*, *Cupressus arizonica* and *Cupressus macrocarpa*); *Juniperus* (e.g. *Juniperus sabinoides*, *Juniperus virginiana*, *Juniperus communis* and *Juniperus ashei*); *Thuya* (e.g. *Thuya orientalis*); *Chamaecyparis* (e.g. *Chamaecyparis obtusa*); *Periplaneta* (e.g. *Periplaneta americana*); *Agropyron* (e.g. *Agropyron repens*); *Secale* (e.g. *Secale cereale*); *Triticum* (e.g. *Triticum aestivum*); *Dactylis* (e.g. *Dactylis glomerata*); *Festuca* (e.g. *Festuca elatior*); *Poa* (e.g. *Poa pratensis* or *Poa compressa*); *Avena* (e.g. *Avena sativa*); *Holcus* (e.g. *Holcus lanatus*); *Anthoxanthum* (e.g. *Anthoxanthum odoratum*); *Arrhenatherum* (e.g. *Arrhenatherum elatius*); *Agrostis* (e.g. *Agrostis alba*); *Phleum* (e.g. *Phleum pratense*); *Phalaris* (e.g. *Phalaris arundinacea*); *Paspalum* (e.g. *Paspalum notatum*); *Sorghum* (e.g. *Sorghum halepensis*); and *Bromus* (e.g. *Bromus inermis*).

- 30       The antigen may be an antigen that is encoded by a nucleic acid vector or it may be not encoded in a nucleic acid vector. In the former case the nucleic acid vector is administered to the subject and the antigen is expressed *in vivo*. In the latter case the antigen may be administered directly to the subject. An antigen not encoded in a nucleic acid vector as used herein refers to any type of antigen that is not a nucleic acid. For instance, in some aspects of the invention the antigen not encoded in a nucleic acid

- 80 -

vector is a polypeptide. Minor modifications of the primary amino acid sequences of polypeptide antigens may also result in a polypeptide which has substantially equivalent antigenic activity as compared to the unmodified counterpart polypeptide. Such modifications may be deliberate, as by site-directed mutagenesis, or may be spontaneous.

5 All of the polypeptides produced by these modifications are included herein as long as antigenicity still exists. The polypeptide may be, for example, a viral polypeptide.

The term substantially purified as used herein refers to a polypeptide which is substantially free of other proteins, lipids, carbohydrates or other materials with which it is naturally associated. One skilled in the art can purify viral or bacterial polypeptides using standard techniques for protein purification. The substantially pure polypeptide  
10 will often yield a single major band on a non-reducing polyacrylamide gel. In the case of partially glycosylated polypeptides or those that have several start codons, there may be several bands on a non-reducing polyacrylamide gel, but these will form a distinctive pattern for that polypeptide. The purity of the viral or bacterial polypeptide can also be  
15 determined by amino-terminal amino acid sequence analysis. Other types of antigens not encoded by a nucleic acid vector such as polysaccharides, small molecule, mimics etc are described above, and included within the invention.

The invention also utilizes polynucleotides encoding the antigenic polypeptides. It is envisioned that the antigen may be delivered to the subject in a nucleic acid  
20 molecule which encodes for the antigen such that the antigen must be expressed *in vivo*. Such antigens delivered to the subject in a nucleic acid vector are referred to as antigens encoded by a nucleic acid vector. The nucleic acid encoding the antigen is operatively linked to a gene expression sequence which directs the expression of the antigen nucleic acid within a eukaryotic cell. The gene expression sequence is any regulatory nucleotide  
25 sequence, such as a promoter sequence or promoter-enhancer combination, which facilitates the efficient transcription and translation of the antigen nucleic acid to which it is operatively linked. The gene expression sequence may, for example, be a mammalian or viral promoter, such as a constitutive or inducible promoter. Constitutive mammalian promoters include, but are not limited to, the promoters for the following genes:  
30 hypoxanthine phosphoribosyl transferase (HPTR), adenosine deaminase, pyruvate kinase, b-actin promoter and other constitutive promoters. Exemplary viral promoters which function constitutively in eukaryotic cells include, for example, promoters from

- 81 -

the cytomegalovirus (CMV), simian virus (e.g., SV40), papilloma virus, adenovirus, human immunodeficiency virus (HIV), Rous sarcoma virus, cytomegalovirus, the long terminal repeats (LTR) of Moloney leukemia virus and other retroviruses, and the thymidine kinase promoter of herpes simplex virus. Other constitutive promoters are known to those of ordinary skill in the art. The promoters useful as gene expression sequences of the invention also include inducible promoters. Inducible promoters are expressed in the presence of an inducing agent. For example, the metallothionein promoter is induced to promote transcription and translation in the presence of certain metal ions. Other inducible promoters are known to those of ordinary skill in the art.

In general, the gene expression sequence shall include, as necessary, 5' non-transcribing and 5' non-translating sequences involved with the initiation of transcription and translation, respectively, such as a TATA box, capping sequence, CAAT sequence, and the like. Especially, such 5' non-transcribing sequences will include a promoter region which includes a promoter sequence for transcriptional control of the operably joined antigen nucleic acid. The gene expression sequences optionally include enhancer sequences or upstream activator sequences as desired.

The antigen nucleic acid is operatively linked to the gene expression sequence. As used herein, the antigen nucleic acid sequence and the gene expression sequence are said to be operably linked when they are covalently linked in such a way as to place the expression or transcription and/or translation of the antigen coding sequence under the influence or control of the gene expression sequence. Two DNA sequences are said to be operably linked if induction of a promoter in the 5' gene expression sequence results in the transcription of the antigen sequence and if the nature of the linkage between the two DNA sequences does not (1) result in the introduction of a frame-shift mutation, (2) interfere with the ability of the promoter region to direct the transcription of the antigen sequence, or (3) interfere with the ability of the corresponding RNA transcript to be translated into a protein. Thus, a gene expression sequence would be operably linked to an antigen nucleic acid sequence if the gene expression sequence were capable of effecting transcription of that antigen nucleic acid sequence such that the resulting transcript is translated into the desired protein or polypeptide.

The antigen nucleic acid of the invention may be delivered to the immune system alone or in association with a vector. In its broadest sense, a vector is any vehicle

capable of facilitating the transfer of the antigen nucleic acid to the cells of the immune system so that the antigen can be expressed and presented on the surface of the immune cell. The vector generally transports the nucleic acid to the immune cells with reduced degradation relative to the extent of degradation that would result in the absence of the vector. The vector optionally includes the above-described gene expression sequence to enhance expression of the antigen nucleic acid in immune cells. In general, the vectors useful in the invention include, but are not limited to, plasmids, phagemids, viruses, other vehicles derived from viral or bacterial sources that have been manipulated by the insertion or incorporation of the antigen nucleic acid sequences. Viral vectors are a preferred type of vector and include, but are not limited to, nucleic acid sequences from the following viruses: retrovirus, such as Moloney murine leukemia virus, Harvey murine sarcoma virus, murine mammary tumor virus, and Rous sarcoma virus; adenovirus, adeno-associated virus; SV40-type viruses; polyoma viruses; Epstein-Barr viruses; papilloma viruses; herpes virus; vaccinia virus; polio virus; and RNA virus such as a retrovirus. One can readily employ other vectors not named but known in the art.

Preferred viral vectors are based on non-cytopathic eukaryotic viruses in which non-essential genes have been replaced with the gene of interest. Non-cytopathic viruses include retroviruses, the life cycle of which involves reverse transcription of genomic viral RNA into DNA with subsequent proviral integration into host cellular DNA. Retroviruses have been approved for human gene therapy trials. Most useful are those retroviruses that are replication-deficient (i.e., capable of directing synthesis of the desired proteins, but incapable of manufacturing an infectious particle). Such genetically altered retroviral expression vectors have general utility for the high-efficiency transduction of genes *in vivo*. Standard protocols for producing replication-deficient retroviruses (including the steps of incorporation of exogenous genetic material into a plasmid, transfection of a packaging cell lined with plasmid, production of recombinant retroviruses by the packaging cell line, collection of viral particles from tissue culture media, and infection of the target cells with viral particles) are provided in Kriegler, M., Gene Transfer and Expression, A Laboratory Manual W.H. Freeman C.O., New York (1990) and Murry, E.J. Methods in Molecular Biology, vol. 7, Humana Press, Inc., Clifton, New Jersey (1991).



- 83 -

A preferred virus for certain applications is the adeno-associated virus, a double-stranded DNA virus. The adeno-associated virus can be engineered to be replication -deficient and is capable of infecting a wide range of cell types and species. It further has advantages such as, heat and lipid solvent stability; high transduction  
5 frequencies in cells of diverse lineages, including hemopoietic cells; and lack of superinfection inhibition thus allowing multiple series of transductions. Reportedly, the adeno-associated virus can integrate into human cellular DNA in a site-specific manner, thereby minimizing the possibility of insertional mutagenesis and variability of inserted gene expression characteristic of retroviral infection. In addition, wild-type  
10 adeno-associated virus infections have been followed in tissue culture for greater than 100 passages in the absence of selective pressure, implying that the adeno-associated virus genomic integration is a relatively stable event. The adeno-associated virus can also function in an extrachromosomal fashion.

Other vectors include plasmid vectors. Plasmid vectors have been extensively  
15 described in the art and are well-known to those of skill in the art. See e.g., Sambrook et al., Molecular Cloning: A Laboratory Manual, Second Edition, Cold Spring Harbor Laboratory Press, 1989. In the last few years, plasmid vectors have been found to be particularly advantageous for delivering genes to cells *in vivo* because of their inability to replicate within and integrate into a host genome. These plasmids, however, having a  
20 promoter compatible with the host cell, can express a peptide from a gene operatively encoded within the plasmid. Some commonly used plasmids include pBR322, pUC18, pUC19, pRC/CMV, SV40, and pBlueScript. Other plasmids are well-known to those of ordinary skill in the art. Additionally, plasmids may be custom designed using restriction enzymes and ligation reactions to remove and add specific fragments of DNA.

25 It has recently been discovered that gene carrying plasmids can be delivered to the immune system using bacteria. Modified forms of bacteria such as *Salmonella* can be transfected with the plasmid and used as delivery vehicles. The bacterial delivery vehicles can be administered to a host subject orally or by other administration means. The bacteria deliver the plasmid to immune cells, e.g. B cells, dendritic cells, likely by  
30 passing through the gut barrier. High levels of immune protection have been established using this methodology. Such methods of delivery are useful for the aspects of the

invention utilizing systemic delivery of antigen, Immunostimulatory nucleic acid and/or other therapeutic agent.

Thus, the immunostimulatory nucleic acids are useful as vaccine adjuvants. It was previously established that CpG oligonucleotides are excellent vaccine adjuvants. It was also demonstrated, however, that CpG ODN which are superb vaccine adjuvants in mice are not the preferred adjuvants in non-rodent animals. In order to identify the best immunostimulatory nucleic acids for use as a vaccine adjuvant in humans and other non-rodent animals, *in vivo* screening of different nucleic acids for this purpose was conducted. Several *in vitro* assays were evaluated in mice for their predictive value of adjuvant activity *in vivo* in mice. During the course of this study, an *in vitro* test that is predictive of *in vivo* efficacy was identified. It was discovered, rather surprisingly, that both B cell and NK cell activation correlated particularly well with the ability of an immunostimulatory nucleic acid to enhance an *in vivo* immune response against an antigen.

The good predictive value of B cell activation for *in vivo* vaccine adjuvant activity is most likely linked to the central role of B cells in the establishment of a specific immune response. Polyclonal proliferation of B cells (induced by immunostimulatory nucleic acids) increases the likelihood of an antigen specific B cell/T helper cell match. Furthermore, enhanced expression of the co-stimulatory molecule CD86 on polyclonally expanded B cells activates antigen specific T helper cells. B cells also increase their CD40 expression in response to immunostimulatory nucleic acids improving the capability of CD40L expressing activated T helper cells to stimulate B cells. Increased ICAM-1 synthesis on B cells facilitates the cell to cell contact. Thus, the activation status of polyclonal B cells plays a critical role during the initiation of a specific antibody response.

The contribution of NK cell activity for the establishment of specific antibodies was, however, surprising. NK cells are part of the innate immune system and as such are involved in the first line of defense against pathogens. Most likely the cytokine pattern produced by NK cells upon activation is closely related to the initiation of a specific immune response. Thus, in one aspect the invention relates to a method of identifying an adjuvant, by detecting NK cell activation. The NK cell activation assay may be carried out as described in the Examples below or using other known NK cell activity assays. It

- 85 -

is preferred, however that a mixed cell population such as PBMC be used because of the likelihood that NK cell activation is an indirect effect. The assay is preferably useful for identifying immunostimulatory nucleic acids which are useful as adjuvants in human and other non-rodent animals.

5 Cytokine induction was also identified as an important predictor of *in vivo* adjuvant activity. As there is a 2 log higher endotoxin sensitivity of human than mouse primary monocytes, some caution, however, is required to avoid endotoxin contamination of immunostimulatory nucleic acids used for testing in the human system (Hartmann G., and Krieg A. M. 1999. *Gene Therapy* 6:893). Since TNF- $\alpha$ , IL-6 and IL-  
10 12 are produced by human monocytes in response to even low amounts of endotoxin, their value for high throughput *in vitro* screening assays is limited. On the other hand, human B cells and NK cells show only minor activation by endotoxin and thus are far more useful in testing for immunostimulatory activity.

Stimulation of cellular function in either NK or B cells (i.e., lytic activity,  
15 proliferation) requires a stronger immunostimulatory nucleic acid than the induction of activation markers at their surface (CD69, CD86). For both cell types, the use of cell surface activation markers showed a higher nonspecific background attributable to the phosphorothioate backbone compared to the functional assays. This high sensitivity of surface markers requires the use of low immunostimulatory nucleic acid concentrations  
20 for optimal discrimination between immunostimulatory nucleic acid of similar activity. Thus, the use of surface markers allows the comparison of immunostimulatory nucleic acids with weak activity, while functional assays are preferred for comparing immunostimulatory nucleic acids with high activity. It is of note that the optimal immunostimulatory nucleic acid concentrations for stimulating B cells and NK cells  
25 differ. While 0.6  $\mu\text{g/ml}$  ODN is already maximal to stimulate B cells, optimal NK cell activation may require 6  $\mu\text{g/ml}$  ODN. Both B cell activation and NK cell functional activity were measured within freshly isolated PBMC. It was previously found that highly purified human primary B cells are activated by CpG DNA. The existence of a direct effect of CpG DNA on NK cells is less clear, and a secondary mechanism  
30 mediated by another cell type within PBMC might contribute to CpG-induced functional activity of NK cells.

The nucleic acids of the invention may be administered to a subject with an anti-microbial agent. An anti-microbial agent, as used herein, refers to a naturally-occurring or synthetic compound which is capable of killing or inhibiting infectious microorganisms. The type of anti-microbial agent useful according to the invention will  
5 depend upon the type of microorganism with which the subject is infected or at risk of becoming infected. Anti-microbial agents include but are not limited to anti-bacterial agents, anti-viral agents, anti-fungal agents and anti-parasitic agents. Phrases such as “anti-infective agent”, “anti-bacterial agent”, “anti-viral agent”, “anti-fungal agent”, “anti-parasitic agent” and “parasiticide” have well-established meanings to those of  
10 ordinary skill in the art and are defined in standard medical texts. Briefly, anti-bacterial agents kill or inhibit bacteria, and include antibiotics as well as other synthetic or natural compounds having similar functions. Antibiotics are low molecular weight molecules which are produced as secondary metabolites by cells, such as microorganisms. In general, antibiotics interfere with one or more bacterial functions or structures which are  
15 specific for the microorganism and which are not present in host cells. Anti-viral agents can be isolated from natural sources or synthesized and are useful for killing or inhibiting viruses. Anti-fungal agents are used to treat superficial fungal infections as well as opportunistic and primary systemic fungal infections. Anti-parasite agents kill or inhibit parasites.

20 Examples of anti-parasitic agents, also referred to as parasiticides useful for human administration include but are not limited to albendazole, amphotericin B, benznidazole, bithionol, chloroquine HCl, chloroquine phosphate, clindamycin, dehydroemetine, diethylcarbamazine, diloxanide furoate, eflornithine, furazolidone, glucocorticoids, halofantrine, iodoquinol, ivermectin, mebendazole, mefloquine,  
25 meglumine antimoniate, melarsoprol, metrifonate, metronidazole, niclosamide, nifurtimox, oxamniquine, paromomycin, pentamidine isethionate, piperazine, praziquantel, primaquine phosphate, proguanil, pyrantel pamoate, pyrimethamine-sulfonamides, pyrimethamine-sulfadoxine, quinacrine HCl, quinine sulfate, quinidine gluconate, spiramycin, stibogluconate sodium (sodium antimony gluconate), suramin,  
30 tetracycline, doxycycline, thiabendazole, tinidazole, trimethoprim-sulfamethoxazole, and tryparsamide some of which are used alone or in combination with others.

Parasiticides used in non-human subjects include piperazine, diethylcarbamazine, thiabendazole, fenbendazole, albendazole, oxfendazole, oxibendazole, febantel, levamisole, pyrantel tartrate, pyrantel pamoate, dichlorvos, ivermectin, doramectin, milbemycin oxime, iprinomectin, moxidectin, N-butyl chloride, 5 toluene, hygromycin B thiacetarsemide sodium, melarsomine, praziquantel, epsiprantel, benzimidazoles such as fenbendazole, albendazole, oxfendazole, clorsulon, albendazole, amprolium; decoquinate, lasalocid, monensin sulfadimethoxine; sulfamethazine, sulfaquinoxaline, metronidazole.

Parasiticides used in horses include mebendazole, oxfendazole, febantel, 10 pyrantel, dichlorvos, trichlorfon, ivermectin, piperazine; for *S. westeri*: ivermectin, benzimidazoles such as thiabendazole, cambendazole, oxibendazole and fenbendazole. Useful parasiticides in dogs include milbemycin oxime, ivermectin, pyrantel pamoate and the combination of ivermectin and pyrantel. The treatment of parasites in swine can include the use of levamisole, piperazine, pyrantel, thiabendazole, dichlorvos and 15 fenbendazole. In sheep and goats anthelmintic agents include levamisole or ivermectin. Caparsolate has shown some efficacy in the treatment of *D. immitis* (heartworm) in cats.

Antibacterial agents kill or inhibit the growth or function of bacteria. A large class of antibacterial agents is antibiotics. Antibiotics, which are effective for killing or inhibiting a wide range of bacteria, are referred to as broad spectrum antibiotics. Other 20 types of antibiotics are predominantly effective against the bacteria of the class gram-positive or gram-negative. These types of antibiotics are referred to as narrow spectrum antibiotics. Other antibiotics which are effective against a single organism or disease and not against other types of bacteria, are referred to as limited spectrum antibiotics. Antibacterial agents are sometimes classified based on their primary mode of action. In 25 general, antibacterial agents are cell wall synthesis inhibitors, cell membrane inhibitors, protein synthesis inhibitors, nucleic acid synthesis or functional inhibitors, and competitive inhibitors.

Anti-bacterial agents useful in the invention include but are not limited to natural penicillins, semi-synthetic penicillins, clavulanic acid, cephalosporins, bacitracin, 30 ampicillin, carbenicillin, oxacillin, azlocillin, mezlocillin, piperacillin, methicillin, dicloxacillin, nafcillin, cephalothin, cephapirin, cephalexin, cefamandole, cefaclor, cefazolin, cefuroxime, cefoxitin, cefotaxime, cefsulodin, cefetamet, cefixime, ceftriaxone,

cefoperazone, ceftazidime, moxalactam, carbapenems, imipenems, monobactams,  
 euztreonam, vancomycin, polymyxin, amphotericin B, nystatin, imidazoles,  
 clotrimazole, miconazole, ketoconazole, itraconazole, fluconazole, rifampins,  
 ethambutol, tetracyclines, chloramphenicol, macrolides, aminoglycosides, streptomycin,  
 5 kanamycin, tobramycin, amikacin, gentamicin, tetracycline, minocycline, doxycycline,  
 chlortetracycline, erythromycin, roxithromycin, clarithromycin, oleandomycin,  
 azithromycin, chloramphenicol, quinolones, co-trimoxazole, norfloxacin, ciprofloxacin,  
 enoxacin, nalidixic acid, temafloxacin, sulfonamides, gantrisin, and trimethoprim;  
 Acedapsone ; Acetosulfone Sodium; Alamecin; Alexidine; Amdinocillin; Amdinocillin  
 10 Pivoxil; Amicycline; Amifloxacin; Amifloxacin Mesylate; Amikacin; Amikacin Sulfate;  
 Aminosalicilic acid; Aminosalicylate sodium; Amoxicillin; Amphomycin; Ampicillin;  
 Ampicillin Sodium; Apalcillin Sodium; Apramycin; Aspartocin; Astromicin Sulfate;  
 Avilamycin; Avoparcin; Azithromycin; Azlocillin; Azlocillin Sodium; Bacampicillin  
 Hydrochloride; Bacitracin; Bacitracin Methylene Disalicylate; Bacitracin Zinc;  
 15 Bambermycins; Benzoylpas Calcium; Berythromycin ; Betamicin Sulfate; Biapenem;  
 Biniramycin; Biphenamine Hydrochloride ; Bispyrithione Magsulfex ; Butikacin;  
 Butirosin Sulfate; Capreomycin Sulfate; Carbadox; Carbenicillin Disodium;  
 Carbenicillin Indanyl Sodium; Carbenicillin Phenyl Sodium; Carbenicillin Potassium;  
 Carumonam Sodium; Cefaclor; Cefadroxil; Cefamandole; Cefamandole Nafate;  
 20 Cefamandole Sodium; Cefaparole; Cefatrizine; Cefazaflur Sodium; Cefazolin; Cefazolin  
 Sodium; Cefbuperazone; Cefdinir; Cefepime; Cefepime Hydrochloride; Cefetecol;  
 Cefixime; Cefmenoxime Hydrochloride; Cefmetazole; Cefmetazole Sodium; Cefonicid  
 Monosodium; Cefonicid Sodium; Cefoperazone Sodium; Ceforanide; Cefotaxime  
 Sodium; Cefotetan; Cefotetan Disodium; Cefotiam Hydrochloride; Cefoxitin; Cefoxitin  
 25 Sodium; Cefpimizole; Cefpimizole Sodium; Cefpiramide; Cefpiramide Sodium;  
 Cefpirome Sulfate; Cefpodoxime Proxetil; Cefprozil; Cefroxadine; Cefsulodin Sodium;  
 Ceftazidime; Ceftibuten; Ceftizoxime Sodium; Ceftriaxone Sodium; Cefuroxime;  
 Cefuroxime Axetil; Cefuroxime Pivoxetil; Cefuroxime Sodium; Cephacetrile Sodium;  
 Cephalexin; Cephalexin Hydrochloride; Cephaloglycin; Cephaloridine; Cephalothin  
 30 Sodium; Cephapirin Sodium; Cephradine; Cetocycline Hydrochloride; Cetophenicol;  
 Chloramphenicol ; Chloramphenicol Palmitate ; Chloramphenicol Pantothenate Complex  
 ; Chloramphenicol Sodium Succinate; Chlorhexidine Phosphanilate; Chloroxylonol;

- Chlortetracycline Bisulfate ; Chlortetracycline Hydrochloride ; Cinoxacin;  
Ciprofloxacin; Ciprofloxacin Hydrochloride; Cirolemycin ; Clarithromycin;  
Clinafloxacin Hydrochloride; Clindamycin; Clindamycin Hydrochloride; Clindamycin  
Palmitate Hydrochloride; Clindamycin Phosphate; Clofazimine ; Cloxacillin Benzathine;  
5 Cloxacillin Sodium; Cloxyquin; Colistimethate Sodium; Colistin Sulfate; Coumermycin;  
Coumermycin Sodium; Cyclacillin; Cycloserine; Dalfopristin; Dapsone ; Daptomycin;  
Demeclocycline; Demeclocycline Hydrochloride; Demecycline; Denofungin ;  
Diaveridine; Dicloxacillin; Dicloxacillin Sodium; Dihydrostreptomycin Sulfate;  
Dipyrrithione; Dirithromycin; Doxycycline; Doxycycline Calcium ; Doxycycline  
10 Fosfatex; Doxycycline Hyclate; Droxacin Sodium; Enoxacin; Epicillin; Epitetracycline  
Hydrochloride; Erythromycin; Erythromycin Acistrate; Erythromycin Estolate;  
Erythromycin Ethylsuccinate; Erythromycin Gluceptate; Erythromycin Lactobionate;  
Erythromycin Propionate; Erythromycin Stearate; Ethambutol Hydrochloride;  
Ethionamide; Fleroxacin; Floxacillin; Fludalanine; Flumequine; Fosfomycin;  
15 Fosfomycin Tromethamine; Fumoxicillin; Furazolium Chloride; Furazolium Tartrate;  
Fusidate Sodium; Fusidic Acid; Gentamicin Sulfate; Gloximonom; Gramicidin;  
Haloprogin; Hetacillin; Hetacillin Potassium; Hexedine; Ibafoxacin; Imipenem;  
Isoconazole; Isepamicin; Isoniazid; Josamycin; Kanamycin Sulfate; Kitasamycin;  
Levofuraltadone; Levopropylcillin Potassium; Lexithromycin; Lincomycin; Lincomycin  
20 Hydrochloride; Lomefloxacin; Lomefloxacin Hydrochloride; Lomefloxacin Mesylate;  
Loracarbef; Mafenide; Meclocycline; Meclocycline Sulfosalicylate; Megalomycin  
Potassium Phosphate; Mequidox; Meropenem; Methacycline; Methacycline  
Hydrochloride; Methenamine; Methenamine Hippurate; Methenamine Mandelate;  
Methicillin Sodium; Metioprime; Metronidazole Hydrochloride; Metronidazole  
25 Phosphate; Mezlocillin; Mezlocillin Sodium; Minocycline; Minocycline Hydrochloride;  
Mirincamycin Hydrochloride ; Monensin ; Monensin Sodium ; Nafcillin Sodium;  
Nalidixate Sodium; Nalidixic Acid; Natamycin; Nebramycin; Neomycin Palmitate;  
Neomycin Sulfate; Neomycin Undecylenate ; Netilmicin Sulfate; Neutramycin;  
Nifuradene; Nifuraldezone; Nifuratel ; Nifuratrone; Nifurdazil; Nifurimide; Nifurpirinol;  
30 Nifurquinazol; Nifurthiazole; Nitrocyline; Nitrofurantoin; Nitromide; Norfloxacin;  
Novobiocin Sodium; Ofloxacin; Ormetoprim; Oxacillin Sodium; Oximonam; Oximonam  
Sodium; Oxolinic Acid; Oxytetracycline; Oxytetracycline Calcium; Oxytetracycline

- Hydrochloride; Paldimycin; Parachlorophenol; Paulomycin; Pefloxacin; Pefloxacin Mesylate; Penamecillin; Penicillin G Benzathine; Penicillin G Potassium; Penicillin G Procaine; Penicillin G Sodium; Penicillin V; Penicillin V Benzathine; Penicillin V Hydrabamine; Penicillin V Potassium; Pentizidone Sodium; Phenyl Aminosalicylate;
- 5 Piperacillin Sodium; Pirbenicillin Sodium; Piridicillin Sodium; Pirlimycin Hydrochloride; Pivampicillin Hydrochloride; Pivampicillin Pamoate; Pivampicillin Probenate; Polymyxin B Sulfate; Porfiromycin ; Propikacin; Pyrazinamide; Pyrrithione Zinc; Quindecamine Acetate; Quinupristin; Racephenicol; Ramoplanin; Ranimycin; Relomycin; Repromicin; Rifabutin; Rifametan; Rifamexil; Rifamide; Rifampin;
- 10 Rifapentine; Rifaximin; Rolitettracycline; Rolitettracycline Nitrate; Rosaramicin; Rosaramicin Butyrate; Rosaramicin Propionate; Rosaramicin Sodium Phosphate; Rosaramicin Stearate; Rosoxacin; Roxarsone; Roxithromycin; Sancycline; Sanfetrinem Sodium; Sarmoxicillin; Sarpicillin; Scopafungin ; Sisomicin; Sisomicin Sulfate; Sparfloxacin; Spectinomycin Hydrochloride; Spiramycin; Stallimycin Hydrochloride;
- 15 Steffimycin; Streptomycin Sulfate; Streptonicozid; Sulfabenz ; Sulfabenzamide; Sulfacetamide; Sulfacetamide Sodium; Sulfacytine; Sulfadiazine; Sulfadiazine Sodium; Sulfadoxine; Sulfalene; Sulfamerazine; Sulfameter; Sulfamethazine; Sulfamethizole; Sulfamethoxazole; Sulfamonomethoxine; Sulfamoxole; Sulfanilate Zinc; Sulfanitran ; Sulfasalazine; Sulfasomizole; Sulfathiazole; Sulfazamet; Sulfisoxazole; Sulfisoxazole
- 20 Acetyl; Sulfisoxazole Diolamine; Sulfomyxin; Sulopenem; Sultamicillin; Suncillin Sodium; Talampicillin Hydrochloride; Teicoplanin; Temafloxacin Hydrochloride; Temocillin; Tetracycline; Tetracycline Hydrochloride; Tetracycline Phosphate Complex; Tetroxoprim; Thiamphenicol; Thiphencillin Potassium; Ticarcillin Cresyl Sodium; Ticarcillin Disodium; Ticarcillin Monosodium; Ticlatone; Tiodonium Chloride;
- 25 Tobramycin; Tobramycin Sulfate; Tosufloxacin; Trimethoprim; Trimethoprim Sulfate; Trisulfapyrimidines; Troleandomycin; Trospectomycin Sulfate; Tyrothricin; Vancomycin; Vancomycin Hydrochloride; Virginiamycin; and Zorbamycin.

Antiviral agents are compounds which prevent infection of cells by viruses or replication of the virus within the cell. There are many fewer antiviral drugs than

30 antibacterial drugs because the process of viral replication is so closely related to DNA replication within the host cell, that non-specific antiviral agents would often be toxic to the host. There are several stages within the process of viral infection which can be



blocked or inhibited by antiviral agents. These stages include, attachment of the virus to the host cell (immunoglobulin or binding peptides), uncoating of the virus (e.g. amantadine), synthesis or translation of viral mRNA (e.g. interferon), replication of viral RNA or DNA (e.g. nucleoside analogues), maturation of new virus proteins (e.g. protease inhibitors), and budding and release of the virus.

Nucleotide analogues are synthetic compounds which are similar to nucleotides, but which have an incomplete or abnormal deoxyribose or ribose group. Once the nucleotide analogues are in the cell, they are phosphorylated, producing the triphosphate formed which competes with normal nucleotides for incorporation into the viral DNA or RNA. Once the triphosphate form of the nucleotide analogue is incorporated into the growing nucleic acid chain, it causes irreversible association with the viral polymerase and thus chain termination. Nucleotide analogues include, but are not limited to, acyclovir (used for the treatment of herpes simplex virus and varicella-zoster virus), gancyclovir (useful for the treatment of cytomegalovirus), idoxuridine, ribavirin (useful for the treatment of respiratory syncytial virus), dideoxyinosine, dideoxycytidine, and zidovudine (azidothymidine).

The interferons are cytokines which are secreted by virus-infected cells as well as immune cells. The interferons function by binding to specific receptors on cells adjacent to the infected cells, causing the change in the cell which protects it from infection by the virus.  $\alpha$  and  $\beta$ -interferon also induce the expression of Class I and Class II MHC molecules on the surface of infected cells, resulting in increased antigen presentation for host immune cell recognition.  $\alpha$  and  $\beta$ -interferons are available as recombinant forms and have been used for the treatment of chronic hepatitis B and C infection. At the dosages which are effective for anti-viral therapy, interferons have severe side effects such as fever, malaise and weight loss.

Immunoglobulin therapy is used for the prevention of viral infection. Immunoglobulin therapy for viral infections is different than bacterial infections, because rather than being antigen-specific, the immunoglobulin therapy functions by binding to extracellular virions and preventing them from attaching to and entering cells which are susceptible to the viral infection. The therapy is useful for the prevention of viral infection for the period of time that the antibodies are present in the host. In general there are two types of immunoglobulin therapies, normal immunoglobulin therapy and

hyper-immunoglobulin therapy. Normal immune globulin therapy utilizes a antibody product which is prepared from the serum of normal blood donors and pooled. This pooled product contains low titers of antibody to a wide range of human viruses, such as hepatitis A, parvovirus, enterovirus (especially in neonates). Hyper-immune globulin  
5 therapy utilizes antibodies which are prepared from the serum of individuals who have high titers of an antibody to a particular virus. Those antibodies are then used against a specific virus. Examples of hyper-immune globulins include zoster immune globulin (useful for the prevention of varicella in immuno-compromised children and neonates), human rabies immunoglobulin (useful in the post-exposure prophylaxis of a subject  
10 bitten by a rabid animal), hepatitis B immune globulin (useful in the prevention of hepatitis B virus, especially in a subject exposed to the virus), and RSV immune globulin (useful in the treatment of respiratory syncytial virus infections).

Another type of immunoglobulin therapy is active immunization. This involves the administration of antibodies or antibody fragments to viral surface proteins. Two  
15 types of vaccines which are available for active immunization of hepatitis B include serum-derived hepatitis B antibodies and recombinant hepatitis B antibodies. Both are prepared from HBsAg. The antibodies are administered in three doses to subjects at high risk of infection with hepatitis B virus, such as health care workers, sexual partners of chronic carriers, and infants.

20 Thus, anti-viral agents useful in the invention include but are not limited to immunoglobulins, amantadine, interferon, nucleoside analogues, and protease inhibitors. Specific examples of anti-virals include but are not limited to Acemannan; Acyclovir; Acyclovir Sodium; Adefovir; Alovudine; Alvircept Sudotox; Amantadine Hydrochloride; Aranotin; Arildone; Ateviridine Mesylate; Avridine; Cidofovir;  
25 Cipamfylline; Cytarabine Hydrochloride; Delavirdine Mesylate; Desciclovir; Didanosine; Disoxaril; Edoxudine; Enviroxime; Famciclovir; Famotone Hydrochloride; Fiacitabine; Fialuridine; Fosarilate; Foscarnet Sodium; Fosfonet Sodium; Ganciclovir; Ganciclovir Sodium; Idoxuridine; Kethoxal; Lamivudine; Lobucavir; Memotone Hydrochloride; Methisazone; Nevirapine; Penciclovir; Pirodavis; Ribavirin;  
30 Rimantadine Hydrochloride; Saquinavir Mesylate; Somantadine Hydrochloride; Sorivudine; Statolon; Stavudine; Tilorone Hydrochloride; Trifluridine; Valacyclovir

- 93 -

Hydrochloride; Vidarabine; Vidarabine Phosphate; Vidarabine Sodium Phosphate; Viroxime; Zalcitabine; Zidovudine; and Ziniviroxime.

Anti-fungal agents are useful for the treatment and prevention of infective fungi.

Anti-fungal agents are sometimes classified by their mechanism of action. Some anti-

- 5 fungal agents function as cell wall inhibitors by inhibiting glucose synthase. These include, but are not limited to, basiungin/ECB. Other anti-fungal agents function by destabilizing membrane integrity. These include, but are not limited to, imidazoles, such as clotrimazole, sertaconazole, fluconazole, itraconazole, ketoconazole, miconazole, and voriconazole, as well as FK 463, amphotericin B, BAY 38-9502, MK 991,
- 10 pradimicin, UK 292, butenafine, and terbinafine. Other anti-fungal agents function by breaking down chitin (e.g. chitinase) or immunosuppression (501 cream). Some examples of commercially-available agents are shown in Table B

**Table B**

Company	Brand Name	Generic Name	Indication	Mechanism of Action
PHARMACIA & UPJOHN	PNU 196443	PNU 196443	Anti Fungal	n/k
Lilly	LY 303366	Basiungin/ECB	Fungal Infections	Anti-fungal/cell wall inhibitor, glucose synthase inhibitor
Bayer	Canesten	Clotrimazole	Fungal Infections	Membrane integrity destabilizer
Fujisawa	FK 463	FK 463	Fungal Infections	Membrane integrity destabilizer
Mylan	Sertaconazole	Sertaconazole	Fungal Infections	Membrane integrity destabilizer
Genzyme	Chitinase	Chitinase	Fungal Infections, Systemic	Chitin Breakdown
Liposome	Abelcet	Amphotericin B, Liposomal	Fungal Infections, Systemic	Membrane integrity destabilizer
Sequus	Amphotec	Amphotericin B, Liposomal	Fungal Infections, Systemic	Membrane integrity destabilizer
Bayer	BAY 38-9502	BAY 38-9502	Fungal Infections, Systemic	Membrane integrity destabilizer
Pfizer	Diflucan	Fluconazole	Fungal Infections, Systemic	Membrane integrity destabilizer
Johnson & Johnson	Sporanox	Itraconazole	Fungal Infections, Systemic	Membrane integrity destabilizer
Sepracor	Itraconazole (2R, 4S)	Itraconazole (2R, 4S)	Fungal Infections, Systemic	Membrane integrity destabilizer
Johnson & Johnson	Nizoral	Ketoconazole	Fungal Infections, Systemic	Membrane integrity destabilizer
Johnson & Johnson	Monistat	Miconazole	Fungal Infections, Systemic	Membrane integrity destabilizer
Merck	MK 991	MK 991	Fungal Infections, Systemic	Membrane integrity destabilizer
Bristol Myers Sq'b	Pradimicin	Pradimicin	Fungal Infections, Systemic	Membrane integrity destabilizer
Pfizer	UK-292, 663	UK-292, 663	Fungal Infections, Systemic	Membrane integrity destabilizer
Pfizer	Voriconazole	Voriconazole	Fungal Infections, Systemic	Membrane integrity destabilizer
Mylan	501 Cream	501 Cream	Inflammatory Fungal Conditions	Immunosuppression
Mylan	Mentax	Butenafine	Nail Fungus	Membrane Integrity Destabiliser
Schering Plough	Anti Fungal	Anti Fungal	Opportunistic Infections	Membrane Integrity Destabiliser
Alza	Mycelex Troche	Clotrimazole	Oral Thrush	Membrane Integrity Stabiliser
Novartis	Lamisil	Terbinafine	Systemic Fungal Infections, Onychomycosis	Membrane Integrity Destabiliser

Thus, the anti-fungal agents useful in the invention include but are not limited to

- 15 imidazoles, FK 463, amphotericin B, BAY 38-9502, MK 991, pradimicin, UK 292, butenafine, chitinase, 501 cream, Acrisorcin; Ambruticin; Amorolfine, Amphotericin B; Azaconazole; Azaserine; Basifungin; Bifonazole; Biphenamine Hydrochloride ; Bispyrithione Magsulfex ; Butoconazole Nitrate; Calcium Undecylenate; Candicidin;

- 94 -

Carbol-Fuchsin; Chlordantoin; Ciclopirox; Ciclopirox Olamine; Cilofungin;  
Cisconazole; Clotrimazole; Cuprimyxin ; Denofungin ; Dipyrithione; Doconazole;  
Econazole; Econazole Nitrate; Enilconazole; Ethonam Nitrate; Fenticonazole Nitrate;  
Filipin; Fluconazole; Flucytosine; Fungimycin; Griseofulvin; Hamycin; Isoconazole ;  
5 Itraconazole; Kalafungin; Ketoconazole; Lomofungin; Lydimycin; Mepartricin ;  
Miconazole; Miconazole Nitrate; Monensin ; Monensin Sodium ; Naftifine  
Hydrochloride; Neomycin Undecylenate ; Nifuratel ; Nifurmerone; Nitralamine  
Hydrochloride; Nystatin; Octanoic Acid; Orconazole Nitrate; Oxiconazole Nitrate;  
Oxifungin Hydrochloride; Parconazole Hydrochloride; Partricin ; Potassium Iodide ;  
10 Proclonol ; Pyrrhithione Zinc ; Pyrrolnitrin; Rutamycin; Sanguinarium Chloride ;  
Saperconazole; Scopafungin ; Selenium Sulfide ; Sinefungin; Sulconazole Nitrate;  
Terbinafine; Terconazole; Thiram; Ticlatone ; Tioconazole; Tolciclate; Tolindate;  
Tolnaftate; Triacetin; Triafungin; Undecylenic Acid; Viridofulvin; Zinc Undecylenate;  
and Zinoconazole Hydrochloride.

15 Immunostimulatory nucleic acids can be combined with other therapeutic agents  
such as adjuvants to enhance immune responses. The immunostimulatory nucleic acid  
and other therapeutic agent may be administered simultaneously or sequentially. When  
the other therapeutic agents are administered simultaneously they can be administered in  
the same or separate formulations, but are administered at the same time. The other  
20 therapeutic agents are administered sequentially with one another and with  
immunostimulatory nucleic acid, when the administration of the other therapeutic agents  
and the immunostimulatory nucleic acid is temporally separated. The separation in time  
between the administration of these compounds may be a matter of minutes or it may be  
longer. Other therapeutic agents include but are not limited to adjuvants, cytokines,  
25 antibodies, antigens, etc.

The immunostimulatory nucleic acids are useful as adjuvants for inducing a  
systemic immune response. Thus either can be delivered to a subject exposed to an  
antigen to produce an enhanced immune response to the antigen.

In addition to the immunostimulatory nucleic acids, the compositions of the  
30 invention may also be administered with non-nucleic acid adjuvants. A non-nucleic acid  
adjuvant is any molecule or compound except for the immunostimulatory nucleic acids  
described herein which can stimulate the humoral and/or cellular immune response.

- 95 -

Non-nucleic acid adjuvants include, for instance, adjuvants that create a depo effect, immune stimulating adjuvants, and adjuvants that create a depo effect and stimulate the immune system.

An adjuvant that creates a depo effect as used herein is an adjuvant that causes the antigen to be slowly released in the body, thus prolonging the exposure of immune cells to the antigen. This class of adjuvants includes but is not limited to alum (e.g., aluminum hydroxide, aluminum phosphate); or emulsion-based formulations including mineral oil, non-mineral oil, water-in-oil or oil-in-water-in oil emulsion, oil-in-water emulsions such as Seppic ISA series of Montanide adjuvants (e.g., Montanide ISA 720, AirLiquide, Paris, France); MF-59 (a squalene-in-water emulsion stabilized with Span 85 and Tween 80; Chiron Corporation, Emeryville, CA; and PROVAX (an oil-in-water emulsion containing a stabilizing detergent and a micelle-forming agent; IDEC, Pharmaceuticals Corporation, San Diego, CA).

An immune stimulating adjuvant is an adjuvant that causes activation of a cell of the immune system. It may, for instance, cause an immune cell to produce and secrete cytokines. This class of adjuvants includes but is not limited to saponins purified from the bark of the *Q. saponaria* tree, such as QS21 (a glycolipid that elutes in the 21<sup>st</sup> peak with HPLC fractionation; Aquila Biopharmaceuticals, Inc., Worcester, MA); poly[di(carboxylatophenoxy)phosphazene (PCPP polymer; Virus Research Institute, USA); derivatives of lipopolysaccharides such as monophosphoryl lipid A (MPL; Ribi ImmunoChem Research, Inc., Hamilton, MT), muramyl dipeptide (MDP; Ribi) and threonyl-muramyl dipeptide (t-MDP; Ribi); OM-174 (a glucosamine disaccharide related to lipid A; OM Pharma SA, Meyrin, Switzerland); and Leishmania elongation factor (a purified *Leishmania* protein; Corixa Corporation, Seattle, WA).

Adjuvants that create a depo effect and stimulate the immune system are those compounds which have both of the above- identified functions. This class of adjuvants includes but is not limited to ISCOMS (Immunostimulating complexes which contain mixed saponins, lipids and form virus-sized particles with pores that can hold antigen; CSL, Melbourne, Australia); SB-AS2 (SmithKline Beecham adjuvant system #2 which is an oil-in-water emulsion containing MPL and QS21; SmithKline Beecham Biologicals [SBB], Rixensart, Belgium); SB-AS4 (SmithKline Beecham adjuvant system #4 which contains alum and MPL; SBB, Belgium); non-ionic block copolymers that form micelles

- 96 -

such as CRL 1005 (these contain a linear chain of hydrophobic polyoxpropylene flanked by chains of polyoxyethylene; Vaxcel, Inc., Norcross, GA); and Syntex Adjuvant Formulation (SAF, an oil-in-water emulsion containing Tween 80 and a nonionic block copolymer; Syntex Chemicals, Inc., Boulder, CO).

5           The immunostimulatory nucleic acids are also useful as mucosal adjuvants. It has previously been discovered that both systemic and mucosal immunity are induced by mucosal delivery of CpG nucleic acids. The systemic immunity induced in response to CpG nucleic acids included both humoral and cell-mediated responses to specific antigens that were not capable of inducing systemic immunity when administered alone  
10 to the mucosa. Furthermore, both CpG nucleic acids and cholera toxin (CT, a mucosal adjuvant that induces a Th2-like response) induced CTL. This was surprising since with systemic immunization, the presence of Th2-like antibodies is normally associated with a lack of CTL (Schirmbeck *et al.*, 1995). Based on the results presented herein it is expected that the immunostimulatory nucleic acids will function in a similar manner.

15           Additionally, the immunostimulatory nucleic acids induce a mucosal response at both local (e.g., lung) and remote (e.g., lower digestive tract) mucosal sites. Significant levels of IgA antibodies are induced at distant mucosal sites by the immunostimulatory nucleic acids. CT is generally considered to be a highly effective mucosal adjuvant. As has been previously reported (Snider 1995), CT induces predominantly IgG1 isotype of  
20 antibodies, which are indicative of Th2-type response. In contrast, the immunostimulatory nucleic acids are more Th1 with predominantly IgG2a antibodies, especially after boost or when the two adjuvants are combined. Th1-type antibodies in general have better neutralizing capabilities, and furthermore, a Th2 response in the lung is highly undesirable because it is associated with asthma (Kay, 1996, Hogg, 1997).

25           Thus the use of immunostimulatory nucleic acids as a mucosal adjuvant has benefits that other mucosal adjuvants cannot achieve. The immunostimulatory nucleic acids of the invention also are useful as mucosal adjuvants for induction of both a systemic and a mucosal immune response.

          Mucosal adjuvants referred to as non-nucleic acid mucosal adjuvants may also be  
30 administered with the Immunostimulatory nucleic acids. A non-nucleic acid mucosal adjuvant as used herein is an adjuvant other than a immunostimulatory nucleic acid that is capable of inducing a mucosal immune response in a subject when administered to a

mucosal surface in conjunction with an antigen. Mucosal adjuvants include but are not limited to Bacterial toxins e.g., Cholera toxin (CT), CT derivatives including but not limited to CT B subunit (CTB) (Wu et al., 1998, Tochikubo et al., 1998); CTD53 (Val to Asp) (Fontana et al., 1995); CTK97 (Val to Lys) (Fontana et al., 1995); CTK104 (Tyr to Lys) (Fontana et al., 1995); CTD53/K63 (Val to Asp, Ser to Lys) (Fontana et al., 1995); CTH54 (Arg to His) (Fontana et al., 1995); CTN107 (His to Asn) (Fontana et al., 1995); CTE114 (Ser to Glu) (Fontana et al., 1995); CTE112K (Glu to Lys) (Yamamoto et al., 1997a); CTS61F (Ser to Phe) (Yamamoto et al., 1997a, 1997b); CTS106 (Pro to Lys) (Douce et al., 1997, Fontana et al., 1995); and CTK63 (Ser to Lys) (Douce et al., 1997, Fontana et al., 1995), Zonula occludens toxin, zot, *Escherichia coli* heat-labile enterotoxin, Labile Toxin (LT), LT derivatives including but not limited to LT B subunit (LTB) (Verweij et al., 1998); LT7K (Arg to Lys) (Komase et al., 1998, Douce et al., 1995); LT61F (Ser to Phe) (Komase et al., 1998); LT112K (Glu to Lys) (Komase et al., 1998); LT118E (Gly to Glu) (Komase et al., 1998); LT146E (Arg to Glu) (Komase et al., 1998); LT192G (Arg to Gly) (Komase et al., 1998); LTK63 (Ser to Lys) (Marchetti et al., 1998, Douce et al., 1997, 1998, Di Tommaso et al., 1996); and LTR72 (Ala to Arg) (Giuliani et al., 1998), Pertussis toxin, PT. (Lycke et al., 1992, Spangler BD, 1992, Freytag and Clemments, 1999, Roberts et al., 1995, Wilson et al., 1995) including PT-9K/129G (Roberts et al., 1995, Cropley et al., 1995); Toxin derivatives (see below) (Holmgren et al., 1993, Verweij et al., 1998, Rappuoli et al., 1995, Freytag and Clements, 1999); Lipid A derivatives (e.g., monophosphoryl lipid A, MPL) (Sasaki et al., 1998, Vancott et al., 1998; Muramyl Dipeptide (MDP) derivatives (Fukushima et al., 1996, Ogawa et al., 1989, Michalek et al., 1983, Morisaki et al., 1983); Bacterial outer membrane proteins (e.g., outer surface protein A (OspA) lipoprotein of *Borrelia burgdorferi*, outer membrane protein of *Neisseria meningitidis*) (Marinero et al., 1999, Van de Verg et al., 1996); Oil-in-water emulsions (e.g., MF59) (Barchfield et al., 1999, Verschoor et al., 1999, O'Hagan, 1998); Aluminum salts (Isaka et al., 1998, 1999); and Saponins (e.g., QS21) Aquila Biopharmaceuticals, Inc., Worcester, MA) (Sasaki et al., 1998, MacNeal et al., 1998), ISCOMS, MF-59 (a squalene-in-water emulsion stabilized with Span 85 and Tween 80; Chiron Corporation, Emeryville, CA); the Seppic ISA series of Montanide adjuvants (e.g., Montanide ISA 720; AirLiquide, Paris, France); PROVAX (an oil-in-water emulsion containing a stabilizing detergent and a micelle-

forming agent; IDEC Pharmaceuticals Corporation, San Diego, CA); Syntex Adjuvant Formulation (SAF; Syntex Chemicals, Inc., Boulder, CO); poly[di(carboxylatophenoxy)phosphazene (PCPP polymer; Virus Research Institute, USA) and Leishmania elongation factor (Corixa Corporation, Seattle, WA).

5 Immune responses can also be induced or augmented by the co-administration or co-linear expression of cytokines (Bueler & Mulligan, 1996; Chow *et al.*, 1997; Geissler *et al.*, 1997; Iwasaki *et al.*, 1997; Kim *et al.*, 1997) or B-7 co-stimulatory molecules (Iwasaki *et al.*, 1997; Tsuji *et al.*, 1997) with the Immunostimulatory nucleic acids. The cytokines can be administered directly with Immunostimulatory nucleic acids or may be  
10 administered in the form of a nucleic acid vector that encodes the cytokine, such that the cytokine can be expressed *in vivo*. In one embodiment, the cytokine is administered in the form of a plasmid expression vector. The term cytokine is used as a generic name for a diverse group of soluble proteins and peptides which act as humoral regulators at nano- to picomolar concentrations and which, either under normal or pathological conditions,  
15 modulate the functional activities of individual cells and tissues. These proteins also mediate interactions between cells directly and regulate processes taking place in the extracellular environment. Examples of cytokines include, but are not limited to IL-1, IL-2, IL-4, IL-5, IL-6, IL-7, IL-10, IL-12, IL-15, IL-18, granulocyte-macrophage colony stimulating factor (GM-CSF), granulocyte colony stimulating factor (G-CSF), interferon-  
20  $\gamma$  ( $\gamma$ -IFN), IFN- $\alpha$ , tumor necrosis factor (TNF), TGF- $\beta$ , FLT-3 ligand, and CD40 ligand.

Cytokines play a role in directing the T cell response. Helper (CD4+) T cells orchestrate the immune response of mammals through production of soluble factors that act on other immune system cells, including other T cells. Most mature CD4+ T helper cells express one of two cytokine profiles: Th1 or Th2. The Th1 subset promotes  
25 delayed-type hypersensitivity, cell-mediated immunity, and immunoglobulin class switching to IgG<sub>2a</sub>. The Th2 subset induces humoral immunity by activating B cells, promoting antibody production, and inducing class switching to IgG<sub>1</sub> and IgE. In some embodiments, it is preferred that the cytokine be a Th1 cytokine.

The nucleic acids are also useful for redirecting an immune response from a Th2  
30 immune response to a Th1 immune response. Redirection of an immune response from a Th2 to a Th1 immune response can be assessed by measuring the levels of cytokines produced in response to the nucleic acid (*e.g.*, by inducing monocytic cells and other



cells to produce Th1 cytokines, including IL-12, IFN- $\gamma$  and GM-CSF). The redirection or rebalance of the immune response from a Th2 to a Th1 response is particularly useful for the treatment or prevention of asthma. For instance, an effective amount for treating asthma can be that amount; useful for redirecting a Th2 type of immune response that is associated with asthma to a Th1 type of response. Th2 cytokines, especially IL-4 and IL-5 are elevated in the airways of asthmatic subjects. These cytokines promote important aspects of the asthmatic inflammatory response, including IgE isotype switching, eosinophil chemotaxis and activation and mast cell growth. Th1 cytokines, especially IFN- $\gamma$  and IL-12, can suppress the formation of Th2 clones and production of Th2 cytokines. The immunostimulatory nucleic acids of the invention cause an increase in Th1 cytokines which helps to rebalance the immune system, preventing or reducing the adverse effects associated with a predominately Th2 immune response.

The nucleic acids are also useful for improving survival, differentiation, activation and maturation of dendritic cells. The immunostimulatory nucleic acids have the unique capability to promote cell survival, differentiation, activation and maturation of dendritic cells. Dendritic precursor cells isolated from blood by immunomagnetic cell sorting develop morphologic and functional characteristics of dendritic cells during a two day incubation with GM-CSF. Without GM-CSF these cells undergo apoptosis. The immunostimulatory nucleic acids are superior to GM-CSF in promoting survival and differentiation of dendritic cells (MHC II expression, cell size, granularity). The immunostimulatory nucleic acids also induce maturation of dendritic cells. Since dendritic cells form the link between the innate and the acquired immune system, by presenting antigens as well as through their expression of pattern recognition receptors which detect microbial molecules like LPS in their local environment, the ability to activate dendritic cells with immunostimulatory nucleic acids supports the use of these immunostimulatory nucleic acid based strategies for *in vivo* and *ex-vivo* immunotherapy against disorders such as cancer and allergic or infectious diseases. The immunostimulatory nucleic acids are also useful for activating and inducing maturation of dendritic cells.

Immunostimulatory nucleic acids also increase natural killer cell lytic activity and antibody dependent cellular cytotoxicity (ADCC). ADCC can be performed using a immunostimulatory nucleic acid in combination with an antibody specific for a cellular

- 100 -

target, such as a cancer cell. When the immunostimulatory nucleic acid is administered to a subject in conjunction with the antibody the subject's immune system is induced to kill the tumor cell. The antibodies useful in the ADCC procedure include antibodies which interact with a cell in the body. Many such antibodies specific for cellular targets  
5 have been described in the art and many are commercially available. Examples of these antibodies are listed below among the list of cancer immunotherapies.

The immunostimulatory nucleic acids may also be administered in conjunction with an anti-cancer therapy. Anti-cancer therapies include cancer medicaments, radiation and surgical procedures. As used herein, a "cancer medicament" refers to a  
10 agent which is administered to a subject for the purpose of treating a cancer. As used herein, "treating cancer" includes preventing the development of a cancer, reducing the symptoms of cancer, and/or inhibiting the growth of an established cancer. In other aspects, the cancer medicament is administered to a subject at risk of developing a cancer for the purpose of reducing the risk of developing the cancer. Various types of  
15 medicaments for the treatment of cancer are described herein. For the purpose of this specification, cancer medicaments are classified as chemotherapeutic agents, immunotherapeutic agents, cancer vaccines, hormone therapy, and biological response modifiers.

As used herein, a "cancer medicament" refers to an agent which is administered  
20 to a subject for the purpose of treating a cancer. As used herein, "treating cancer" includes preventing the development of a cancer, reducing the symptoms of cancer, and/or inhibiting the growth of an established cancer. In other aspects, the cancer medicament is administered to a subject at risk of developing a cancer for the purpose of reducing the risk of developing the cancer. Various types of medicaments for the  
25 treatment of cancer are described herein. For the purpose of this specification, cancer medicaments are classified as chemotherapeutic agents, immunotherapeutic agents, cancer vaccines, hormone therapy, and biological response modifiers. Additionally, the methods of the invention are intended to embrace the use of more than one cancer medicament along with the immunostimulatory nucleic acids. As an example, where  
30 appropriate, the immunostimulatory nucleic acids may be administered with a both a chemotherapeutic agent and an immunotherapeutic agent. Alternatively, the cancer medicament may embrace an immunotherapeutic agent and a cancer vaccine, or a

chemotherapeutic agent and a cancer vaccine, or a chemotherapeutic agent, an immunotherapeutic agent and a cancer vaccine all administered to one subject for the purpose of treating a subject having a cancer or at risk of developing a cancer.

Cancer medicaments function in a variety of ways. Some cancer medicaments work by targeting physiological mechanisms that are specific to tumor cells. Examples include the targeting of specific genes and their gene products (i.e., proteins primarily) which are mutated in cancers. Such genes include but are not limited to oncogenes (e.g., Ras, Her2, bcl-2), tumor suppressor genes (e.g., EGF, p53, Rb), and cell cycle targets (e.g., CDK4, p21, telomerase). Cancer medicaments can alternately target signal transduction pathways and molecular mechanisms which are altered in cancer cells. Targeting of cancer cells via the epitopes expressed on their cell surface is accomplished through the use of monoclonal antibodies. This latter type of cancer medicament is generally referred to herein as immunotherapy.

Other cancer medicaments target cells other than cancer cells. For example, some medicaments prime the immune system to attack tumor cells (i.e., cancer vaccines). Still other medicaments, called angiogenesis inhibitors, function by attacking the blood supply of solid tumors. Since the most malignant cancers are able to metastasize (i.e., exist the primary tumor site and seed a distal tissue, thereby forming a secondary tumor), medicaments that impede this metastasis are also useful in the treatment of cancer. Angiogenic mediators include basic FGF, VEGF, angiopoietins, angiostatin, endostatin, TNF $\alpha$ , TNP-470, thrombospondin-1, platelet factor 4, CAI, and certain members of the integrin family of proteins. One category of this type of medicament is a metalloproteinase inhibitor, which inhibits the enzymes used by the cancer cells to exist the primary tumor site and extravasate into another tissue.

Some cancer cells are antigenic and thus can be targeted by the immune system. In one aspect, the combined administration of immunostimulatory nucleic acids and cancer medicaments, particularly those which are classified as cancer immunotherapies, is useful for stimulating a specific immune response against a cancer antigen. A "cancer antigen" as used herein is a compound, such as a peptide, associated with a tumor or cancer cell surface and which is capable of provoking an immune response when expressed on the surface of an antigen presenting cell in the context of an MHC molecule. Cancer antigens, such as those present in cancer vaccines or those used to

- 102 -

prepare cancer immunotherapies, can be prepared from crude cancer cell extracts, as described in Cohen, et al., 1994, *Cancer Research*, 54:1055, or by partially purifying the antigens, using recombinant technology, or de novo synthesis of known antigens.

Cancer antigens can be used in the form of immunogenic portions of a particular antigen or in some instances a whole cell or a tumor mass can be used as the antigen. Such antigens can be isolated or prepared recombinantly or by any other means known in the art.

The theory of immune surveillance is that a prime function of the immune system is to detect and eliminate neoplastic cells before a tumor forms. A basic principle of this theory is that cancer cells are antigenically different from normal cells and thus elicit immune reactions that are similar to those that cause rejection of immunologically incompatible allografts. Studies have confirmed that tumor cells differ, either qualitatively or quantitatively, in their expression of antigens. For example, "tumor-specific antigens" are antigens that are specifically associated with tumor cells but not normal cells. Examples of tumor specific antigens are viral antigens in tumors induced by DNA or RNA viruses. "Tumor-associated" antigens are present in both tumor cells and normal cells but are present in a different quantity or a different form in tumor cells. Examples of such antigens are oncofetal antigens (e.g., carcinoembryonic antigen), differentiation antigens (e.g., T and Tn antigens), and oncogene products (e.g., HER/neu).

Different types of cells that can kill tumor targets *in vitro* and *in vivo* have been identified: natural killer cells (NK cells), cytolytic T lymphocytes (CTLs), lymphokine-activated killer cells (LAKs), and activated macrophages. NK cells can kill tumor cells without having been previously sensitized to specific antigens, and the activity does not require the presence of class I antigens encoded by the major histocompatibility complex (MHC) on target cells. NK cells are thought to participate in the control of nascent tumors and in the control of metastatic growth. In contrast to NK cells, CTLs can kill tumor cells only after they have been sensitized to tumor antigens and when the target antigen is expressed on the tumor cells that also express MHC class I. CTLs are thought to be effector cells in the rejection of transplanted tumors and of tumors caused by DNA viruses. LAK cells are a subset of null lymphocytes distinct from the NK and CTL populations. Activated macrophages can kill tumor cells in a manner that is not antigen

- 103 -

dependent nor MHC restricted once activated. Activated macrophages are through to decrease the growth rate of the tumors they infiltrate. *In vitro* assays have identified other immune mechanisms such as antibody-dependent, cell-mediated cytotoxic reactions and lysis by antibody plus complement. However, these immune effector mechanisms are thought to be less important *in vivo* than the function of NK, CTLs, LAK, and macrophages *in vivo* (for review see Piessens, W.F., and David, J., "Tumor Immunology", In: Scientific American Medicine, Vol. 2, Scientific American Books, N.Y., pp. 1-13, 1996.

The goal of immunotherapy is to augment a patient's immune response to an established tumor. One method of immunotherapy includes the use of adjuvants. Adjuvant substances derived from microorganisms, such as bacillus Calmette-Guerin, heighten the immune response and enhance resistance to tumors in animals.

Immunotherapeutic agents are medicaments which derive from antibodies or antibody fragments which specifically bind or recognize a cancer antigen. As used herein a cancer antigen is broadly defined as an antigen expressed by a cancer cell. Preferably, the antigen is expressed at the cell surface of the cancer cell. Even more preferably, the antigen is one which is not expressed by normal cells, or at least not expressed to the same level as in cancer cells. Antibody-based immunotherapies may function by binding to the cell surface of a cancer cell and thereby stimulate the endogenous immune system to attack the cancer cell. Another way in which antibody-based therapy functions is as a delivery system for the specific targeting of toxic substances to cancer cells. Antibodies are usually conjugated to toxins such as ricin (e.g., from castor beans), calicheamicin and maytansinoids, to radioactive isotopes such as Iodine-131 and Yttrium-90, to chemotherapeutic agents (as described herein), or to biological response modifiers. In this way, the toxic substances can be concentrated in the region of the cancer and non-specific toxicity to normal cells can be minimized. In addition to the use of antibodies which are specific for cancer antigens, antibodies which bind to vasculature, such as those which bind to endothelial cells, are also useful in the invention. This is because generally solid tumors are dependent upon newly formed blood vessels to survive, and thus most tumors are capable of recruiting and stimulating the growth of new blood vessels. As a result, one strategy of many cancer medicaments

is to attack the blood vessels feeding a tumor and/or the connective tissues (or stroma) supporting such blood vessels.

The use of immunostimulatory nucleic acids in conjunction with immunotherapeutic agents such as monoclonal antibodies is able to increase long-term survival through a number of mechanisms including significant enhancement of ADCC (as discussed above), activation of natural killer (NK) cells and an increase in IFN $\alpha$  levels. The nucleic acids when used in combination with monoclonal antibodies serve to reduce the dose of the antibody required to achieve a biological result.

Examples of cancer immunotherapies which are currently being used or which are in development are listed in Table C.

**Table C**

<i>Cancer Immunotherapies in Development or on the Market</i>		
<b>MARKETER</b>	<b>BRAND NAME (GENERIC NAME)</b>	<b>INDICATION</b>
IDEC/Genentech, Inc./Hoffmann-LaRoche (first monoclonal antibody licensed for the treatment of cancer in the U.S.)	Rituxan™ (rituximab, Mabthera) (IDEC-C2B8, chimeric murine/human anti-CD20 MAb)	non-Hodgkin's lymphoma
Genentech/Hoffmann-La Roche	Herceptin, anti-Her2 hMAb	Breast/ovarian
Cytogen Corp.	Quadramet (CYT-424) radiotherapeutic agent	Bone metastases
Centocor/Glaxo/Ajinomoto	Panorex® (17-1A) (murine monoclonal antibody)	Adjuvant therapy for colorectal (Dukes-C)
Centocor/Ajinomoto	Panorex® (17-1A) (chimeric murine monoclonal antibody)	Pancreatic, lung, breast, ovary
IDEC	IDEC-Y2B8 (murine, anti-CD20 MAb labeled with Yttrium-90)	non-Hodgkin's lymphoma
ImClone Systems	BEC2 (anti-idiotypic MAb, mimics the GD <sub>3</sub> epitope) (with BCG)	Small cell lung
ImClone Systems	C225 (chimeric monoclonal antibody to epidermal growth factor receptor (EGFr))	Renal cell
Techniclone International/Alpha Therapeutics	Oncolym (Lym-1 monoclonal antibody linked to 131 iodine)	non-Hodgkin's lymphoma
Protein Design Labs	SMART M195 Ab, humanized	Acute myleoid leukemia
Techniclone Corporation/Cambridge Antibody Technology	<sup>125</sup> I LYM-1 (Oncolym™)	non-Hodgkin's lymphoma
Aronex Pharmaceuticals, Inc.	ATRAGEN®	Acute promyelocytic leukemia
ImClone Systems	C225 (chimeric anti-EGFr monoclonal antibody) + cisplatin or radiation	Head & neck, non-small cell lung cancer

Altarex, Canada	Ovarex (B43.13, anti-idiotypic CA125, mouse MAb)	Ovarian
Coulter Pharma (Clinical results have been positive, but the drug has been associated with significant bone marrow toxicity)	Bexxar (anti-CD20 Mab labeled with <sup>131</sup> I)	non-Hodgkin's lymphoma
Aronex Pharmaceuticals, Inc.	ATRAGEN®	Kaposi's sarcoma
IDEC Pharmaceuticals Corp./Genentech	Rituxan™ (MAb against CD20) pan-B Ab in combo. with chemotherapy	B cell lymphoma
LeukoSite/Ilex Oncology	LDP-03, huMAb to the leukocyte antigen CAMPATH	Chronic lymphocytic leukemia (CLL)
Center of Molecular Immunology	ior t6 (anti CD6, murine MAb) CTCL	Cancer
Medarex/Novartis	MDX-210 (humanized anti-HER-2 bispecific antibody)	Breast, ovarian
Medarex/Novartis	MDX-210 (humanized anti-HER-2 bispecific antibody)	Prostate, non-small cell lung, pancreatic, breast
Medarex	MDX-11 (complement activating receptor (CAR) monoclonal antibody)	Acute myelogenous leukemia (AML)
Medarex/Novartis	MDX-210 (humanized anti-HER-2 bispecific antibody)	Renal and colon
Medarex	MDX-11 (complement activating receptor (CAR) monoclonal antibody)	Ex vivo bone marrow purging in acute myelogenous leukemia (AML)
Medarex	MDX-22 (humanized bispecific antibody, MAb-conjugates) (complement cascade activators)	Acute myeloid leukemia
Cytogen	OV103 (Yttrium-90 labelled antibody)	Ovarian
Cytogen	OV103 (Yttrium-90 labelled antibody)	Prostate
Aronex Pharmaceuticals, Inc.	ATRAGEN®	non-Hodgkin's lymphoma
Glaxo Wellcome plc	3622W94 MAb that binds to EGP40 (17-1A) pancarcinoma antigen on adenocarcinomas	non-small cell lung, prostate (adjuvant)
Genentech	Anti-VEGF, RhuMAb (inhibits angiogenesis)	Lung, breast, prostate, colorectal
Protein Design Labs	Zenapax (SMART Anti-Tac (IL-2 receptor) Ab, humanized)	Leukemia, lymphoma
Protein Design Labs	SMART M195 Ab, humanized	Acute promyelocytic leukemia
ImClone Systems	C225 (chimeric anti-EGFr monoclonal antibody) + taxol	Breast
ImClone Systems (licensed from RPR)	C225 (chimeric anti-EGFr monoclonal antibody) + doxorubicin	prostate
ImClone Systems	C225 (chimeric anti-EGFr monoclonal antibody) + adriamycin	prostate
ImClone Systems	BEC2 (anti-idiotypic MAb, mimics the GD <sub>3</sub> epitope)	Melanoma
Medarex	MDX-210 (humanized anti-HER-2 bispecific	Cancer

	antibody)	
Medarex	MDX-220 (bispecific for tumors that express TAG-72)	Lung, colon, prostate, ovarian, endometrial, pancreatic and gastric
Medarex/Novartis	MDX-210 (humanized anti-HER-2 bispecific antibody)	Prostate
Medarex/Merck KgaA	MDX-447 (humanized anti-EGF receptor bispecific antibody)	EGF receptor cancers (head & neck, prostate, lung, bladder, cervical, ovarian)
Medarex/Novartis	MDX-210 (humanized anti-HER-2 bispecific antibody)	Comb. Therapy with G-CSF for various cancers, esp. breast
IDEC	MELIMMUNE-2 (murine monoclonal antibody therapeutic vaccine )	Melanoma
IDEC	MELIMMUNE-1 (murine monoclonal antibody therapeutic vaccine )	Melanoma
Immunomedics, Inc.	CEACIDE™ (I-131)	Colorectal and other
NeoRx	Pretarget™ radioactive antibodies	non-Hodgkin's B cell lymphoma
Novopharm Biotech, Inc.	NovoMAb-G2 (pancarcinoma specific Ab)	Cancer
Techniclone Corporation/ Cambridge Antibody Technology	TNT (chimeric MAb to histone antigens)	Brain
Techniclone International/ Cambridge Antibody Technology	TNT (chimeric MAb to histone antigens)	Brain
Novopharm	Gliomab-H (Monoclonals - Humanized Abs)	Brain, melanomas, neuroblastomas
Genetics Institute/AHP	GNI-250 Mab	Colorectal
Merck KgaA	EMD-72000 (chimeric-EGF antagonist)	Cancer
Immunomedics	LymphoCide (humanized LL2 antibody)	non-Hodgkin's B-cell lymphoma
Immunex/AHP	CMA 676 (monoclonal antibody conjugate)	Acute myelogenous leukemia
Novopharm Biotech, Inc.	Monopharm-C	Colon, lung, pancreatic
Novopharm Biotech, Inc.	4B5 anti-idiotypic Ab	Melanoma, small-cell lung
Center of Molecular Immunology	ior egf/r3 (anti EGF-R humanized Ab)	Radioimmunotherapy
Center of Molecular Immunology	ior c5 (murine MAb colorectal) for radioimmunotherapy	Colorectal
Creative BioMolecules/ Chiron	BABS (biosynthetic antibody binding site) Proteins	Breast cancer
ImClone Systems/Chugai	FLK-2 (monoclonal antibody to fetal liver kinase-2 (FLK-2))	Tumor-associated angiogenesis
ImmunoGen, Inc.	Humanized MAb/small-drug conjugate	Small-cell lung
Medarex, Inc.	MDX-260 bispecific, targets GD-2	Melanoma, glioma,



		neuroblastoma
Procyon Biopharma, Inc.	ANA Ab	Cancer
Protein Design Labs	SMART ID10 Ab	B-cell lymphoma
Protein Design Labs/Novartis	SMART ABL 364 Ab	Breast, lung, colon
Immunomedics, Inc.	ImmuRAIT-CEA	Colorectal

Yet other types of chemotherapeutic agents which can be used according to the invention include Aminoglutethimide, Asparaginase, Busulfan, Carboplatin, Chlorombucil, Cytarabine HCl, Dactinomycin, Daunorubicin HCl, Estramustine phosphate sodium, Etoposide (VP16-213), Floxuridine, Fluorouracil (5-FU), Flutamide, Hydroxyurea (hydroxycarbamide), Ifosfamide, Interferon Alfa-2a, Alfa-2b, Leuprolide acetate (LHRH-releasing factor analogue), Lomustine (CCNU), Mechlorethamine HCl (nitrogen mustard), Mercaptopurine, Mesna, Mitotane (o.p'-DDD), Mitoxantrone HCl, Octreotide, Plicamycin, Procarbazine HCl, Streptozocin, Tamoxifen citrate, Thioguanine, Thiotepa, Vinblastine sulfate, Amsacrine (m-AMSA), Azacitidine, Erthropoietin, Hexamethylmelamine (HMM), Interleukin 2, Mitoguazone (methyl-GAG; methyl glyoxal bis-guanylhydrazine; MGBG), Pentostatin (2'-deoxycoformycin), Semustine (methyl-CCNU), Teniposide (VM-26) and Vindesine sulfate.

Cancer vaccines are medicaments which are intended to stimulate an endogenous immune response against cancer cells. Currently produced vaccines predominantly activate the humoral immune system (i.e., the antibody dependent immune response). Other vaccines currently in development are focused on activating the cell-mediated immune system including cytotoxic T lymphocytes which are capable of killing tumor cells. Cancer vaccines generally enhance the presentation of cancer antigens to both antigen presenting cells (e.g., macrophages and dendritic cells) and/or to other immune cells such as T cells, B cells, and NK cells.

Although cancer vaccines may take one of several forms, as discussed infra, their purpose is to deliver cancer antigens and/or cancer associated antigens to antigen presenting cells (APC) in order to facilitate the endogenous processing of such antigens by APC and the ultimate presentation of antigen presentation on the cell surface in the context of MHC class I molecules. One form of cancer vaccine is a whole cell vaccine which is a preparation of cancer cells which have been removed from a subject, treated ex vivo and then reintroduced as whole cells in the subject. Lysates of tumor cells can

- 108 -

also be used as cancer vaccines to elicit an immune response. Another form cancer vaccine is a peptide vaccine which uses cancer-specific or cancer-associated small proteins to activate T cells. Cancer-associated proteins are proteins which are not exclusively expressed by cancer cells (i.e., other normal cells may still express these antigens). However, the expression of cancer-associated antigens is generally consistently upregulated with cancers of a particular type. Yet another form of cancer vaccine is a dendritic cell vaccine which includes whole dendritic cells which have been exposed to a cancer antigen or a cancer-associated antigen in vitro. Lysates or membrane fractions of dendritic cells may also be used as cancer vaccines. Dendritic cell vaccines are able to activate antigen-presenting cells directly. Other cancer vaccines include ganglioside vaccines, heat-shock protein vaccines, viral and bacterial vaccines, and nucleic acid vaccines.

The use of immunostimulatory nucleic acids in conjunction with cancer vaccines provides an improved antigen-specific humoral and cell mediated immune response, in addition to activating NK cells and endogenous dendritic cells, and increasing IFN $\alpha$  levels. This enhancement allows a vaccine with a reduced antigen dose to be used to achieve the same beneficial effect. In some instances, cancer vaccines may be used along with adjuvants, such as those described above.

As used herein, the terms "cancer antigen" and "tumor antigen" are used interchangeably to refer to antigens which are differentially expressed by cancer cells and can thereby be exploited in order to target cancer cells. Cancer antigens are antigens which can potentially stimulate apparently tumor-specific immune responses. Some of these antigens are encoded, although not necessarily expressed, by normal cells. These antigens can be characterized as those which are normally silent (i.e., not expressed) in normal cells, those that are expressed only at certain stages of differentiation and those that are temporally expressed such as embryonic and fetal antigens. Other cancer antigens are encoded by mutant cellular genes, such as oncogenes (e.g., activated ras oncogene), suppressor genes (e.g., mutant p53), fusion proteins resulting from internal deletions or chromosomal translocations. Still other cancer antigens can be encoded by viral genes such as those carried on RNA and DNA tumor viruses.

Other vaccines take the form of dendritic cells which have been exposed to cancer antigens in vitro, have processed the antigens and are able to express the cancer

- 109 -

antigens at their cell surface in the context of MHC molecules for effective antigen presentation to other immune system cells.

The immunostimulatory nucleic acids are used in one aspect of the invention in conjunction with cancer vaccines which are dendritic cell based. A dendritic cell is a professional antigen presenting cell. Dendritic cells form the link between the innate and the acquired immune system by presenting antigens and through their expression of pattern recognition receptors which detect microbial molecules like LPS in their local environment. Dendritic cells efficiently internalize, process, and present soluble specific antigen to which it is exposed. The process of internalizing and presenting antigen causes rapid upregulation of the expression of major histocompatibility complex (MHC) and costimulatory molecules, the production of cytokines, and migration toward lymphatic organs where they are believed to be involved in the activation of T cells.

Table D lists a variety of cancer vaccines which are either currently being used or are in development.

**Table D**

<i>Cancer Vaccines in Development or on the Market</i>		
<b>MARKETER</b>	<b>BRAND NAME (GENERIC NAME)</b>	<b>INDICATION</b>
Center of Molecular Immunology	EGF	Cancer
Center of Molecular Immunology		Ganglioside cancer vaccine
Center of Molecular Immunology	Anti-idiotypic	Cancer vaccine
ImClone Systems/Memorial Sloan-Kettering Cancer Center	Gp75 antigen	Melanoma
ImClone Systems/Memorial Sloan-Kettering Cancer Center	Anti-idiotypic Abs	Cancer vaccines
Progenics Pharmaceuticals, Inc.	GMK melanoma vaccine	Melanoma
Progenics Pharmaceuticals, Inc.	MGV ganglioside conjugate vaccine	Lymphoma, colorectal, lung
Corixa	Her2/neu	Breast, ovarian
AltaRex	Ovarex	Ovarian
AVAX Technologies Inc.	M-Vax, autologous whole cell	Melanoma
AVAX Technologies Inc.	O-Vax, autologous whole cell	Ovarian
AVAX Technologies Inc.	L-Vax, autologous whole cell	Leukemia-AML
Biomira Inc./Chiron	Theratope, STn-KLH	Breast, Colorectal

Biomira Inc.	BLP25, MUC-1 peptide vaccine encapsulated in liposomal delivery system	Lung
Biomira Inc.	BLP25, MUC-1 peptide vaccine encapsulated in liposomal delivery system + Liposomal IL-2	Lung
Biomira Inc.	Liposomal idiotypic vaccine	Lymphoma B-cell malignancies
Ribi Immunochem	Melacine, cell lysate	Melanoma
Corixa	Peptide antigens, microsphere delivery sysem and LeIF adjuvant	Breast
Corixa	Peptide antigens, microsphere delivery sysem and LeIF adjuvant	Prostate
Corixa	Peptide antigens, microsphere delivery sysem and LeIF adjuvant	Ovarian
Corixa	Peptide antigens, microsphere delivery sysem and LeIF adjuvant	Lymphoma
Corixa	Peptide antigens, microsphere delivery sysem and LeIF adjuvant	Lung
Virus Research Institute	Toxin/antigen recombinant delivery system	All cancers
Apollon Inc.	Genevax-TCR	T-cell lymphoma
Bavarian Nordic Research Institute A/S	MVA-based (vaccinia virus) vaccine	Melanoma
BioChem Pharma/BioChem Vaccine	PACIS, BCG vaccine	Bladder
Cantab Pharmaceuticals	TA-HPV	Cervical
Cantab Pharmaceuticals	TA-CIN	Cervical
Cantab Pharmaceuticals	DISC-Virus, immunotherapy	Cancer
Pasteur Merieux Connaught	ImmuCyst®/TheraCys® - BCG Immunotherapeutic (Bacillus Calmette-Guerin/Connaught), for intravesical treatment of superficial bladder cancer	Bladder

As used herein, chemotherapeutic agents embrace all other forms of cancer medicaments which do not fall into the categories of immunotherapeutic agents or cancer vaccines. Chemotherapeutic agents as used herein encompass both chemical and biological agents. These agents function to inhibit a cellular activity which the cancer cell is dependent upon for continued survival. Categories of chemotherapeutic agents include alkylating/alkaloid agents, antimetabolites, hormones or hormone analogs, and miscellaneous antineoplastic drugs. Most if not all of these agents are directly toxic to cancer cells and do not require immune stimulation. Combination chemotherapy and

immunostimulatory nucleic acid administration increases the maximum tolerable dose of chemotherapy.

Chemotherapeutic agents which are currently in development or in use in a clinical setting are shown in Table E.

5

**Table E**

<b><i>Cancer Drugs in Development or on the Market</i></b>			
<b>Marketer</b>	<b>Brand Name</b>	<b>Generic Name</b>	<b>Indication</b>
Abbott	TNP 470/AGM 1470	Fragylne	Anti-Angiogenesis in Cancer
Takeda	TNP 470/AGM 1470	Fragylne	Anti-Angiogenesis in Cancer
Scotia	Meglamine GLA	Meglamine GLA	Bladder Cancer
Medeva	Valstar	Valrubicin	Bladder Cancer - Refractory in situ carcinoma
Medeva	Valstar	Valrubicin	Bladder Cancer - Papillary Cancer
Rhone Poulenc	Gliadel Wafer	Carmustaine + Polifepr Osan	Brain Tumor
Warner Lambert	Undisclosed Cancer (b)	Undisclosed Cancer (b)	Cancer
Bristol Myers Squib	RAS Famesyl Transferase Inhibitor	RAS Famesyl Transferase Inhibitor	Cancer
Novartis	MMI 270	MMI 270	Cancer
Bayer	BAY 12-9566	BAY 12-9566	Cancer
Merck	Famesyl Transferase Inhibitor	Famesyl Transferase Inhibitor	Cancer (Solid tumors - pancreas, colon, lung, breast)
Pfizer	PFE	MMP	Cancer, angiogenesis
Pfizer	PFE	Tyrosine Kinase	Cancer, angiogenesis
Lilly	MTA/LY 231514	MTA/LY 231514	Cancer Solid Tumors
Lilly	LY 264618/Lometexol	Lometexol	Cancer Solid Tumors
Scotia	Glamolec	LiGLA (lithium-gamma linolenate)	Cancer, pancreatic, breast, colon
Warner Lambert	CI-994	CI-994	Cancer, Solid Tumors / Leukemia
Schering AG	Angiogenesis inhibitor	Angiogenesis Inhibitor	Cancer / Cardio
Takeda	TNP-470	n/k	Malignant Tumor
Smithkline Beecham	Hycamtin	Topotecan	Metastatic Ovarian Cancer
Novartis	PKC 412	PKC 412	Multi-Drug Resistant Cancer
Novartis	Valspodar	PSC 833	Myeloid Leukemia/Ovarian Cancer
Immunex	Novantrone	Mitoxantrone	Pain related to hormone refractory prostate cancer.
Warner Lambert	Metaret	Suramin	Prostate
Genentech	Anti-VEGF	Anti-VEGF	Prostate / Breast / Colorectal / NSCL Cancer
British Biotech	Batimastat	Batimastat (BB94)	Pterygium
Eisai	E 7070	E 7070	Solid Tumors
Biochem Pharma	BCH-4556	BCH-4556	Solid Tumors
Sankyo	CS-682	CS-682	Solid Tumors
Agouron	AG2037	AG2037	Solid Tumors
IDEC Pharma	9-AC	9-AC	Solid Tumors
Agouron	VEGF/b-FGF Inhibitors	VEGF/b-FGF Inhibitors	Solid Tumors
Agouron	AG3340	AG3340	Solid Tumors / Macular

- 112 -

			Degen
Vertex	Incel	VX-710	Solid Tumors - IV
Vertex	VX-853	VX-853	Solid Tumors - Oral
Zeneca	ZD 0101 (inj)	ZD 0101	Solid Tumors
Novartis	ISI 641	ISI 641	Solid Tumors
Novartis	ODN 698	ODN 698	Solid Tumors
Tanabe Seiyaku	TA 2516	Marimistat	Solid Tumors
British Biotech	Marimastat	Marimastat (BB 2516)	Solid Tumors
Celltech	CDP 845	Aggrecanase Inhibitor	Solid Tumors / Breast Cancer
Chiroscience	D2163	D2163	Solid Tumors / Metastases
Warner Lambert	PD 183805	PD 183805	
Daiichi	DX8951f	DX8951f	Anti-Cancer
Daiichi	Lemonal DP 2202	Lemonal DP 2202	Anti-Cancer
Fujisawa	FK 317	FK 317	Anticancer Antibiotic
Chugai	Picibanil	OK-432	Antimalignant Tumor
Nycomed Amersham	AD 32/valrubicin	Valrubicin	Bladder Cancer-Refractory Insitu Carcinoma
Nycomed Amersham	Metastron	Strontium Derivative	Bone Cancer (adjunct therapy, Pain)
Schering Plough	Temodal	Temozolomide	Brain Tumours
Schering Plough	Temodal	Temozolomide	Brain Tumours
Liposome	Evacet	Doxorubicin, Liposomal	Breast Cancer
Nycomed Amersham	Yewtaxan	Paclitaxel	Breast Cancer Advanced, Ovarian Cancer Advanced
Bristol Myers Squibb	Taxol	Paclitaxel	Breast Cancer Advanced, Ovarian Cancer Advanced, NSCLC
Roche	Xeloda	Capecitabine	Breast Cancer, Colorectal Cancer
Roche	Furtulon	Doxifluridine	Breast Cancer, Colorectal Cancer, Gastric Cancer
Pharmacia & Upjohn	Adriamycin	Doxorubicin	Breast Cancer, Leukemia
Ivax	Cyclopax	Paclitaxel, Oral	Breast/Ovarian Cancer
Rhone Poulenc	Oral Taxoid	Oral Taxoid	Broad Cancer
AHP	Novantrone	Mitoxantrone	Cancer
Sequus	SPI-077	Cisplatin, Stealth	Cancer
Hoechst	HMR 1275	Flavopiridol	Cancer
Pfizer	CP-358, 774	EGFR	Cancer
Pfizer	CP-609, 754	RAS Oncogene Inhibitor	Cancer
Bristol Myers Squibb	BMS-182751	Oral Platinum	Cancer (Lung, Ovarian)
Bristol Myers Squibb	UFT (Tegafur/Uracil)	UFT (Tegafur/Uracil)	Cancer Oral
Johnson & Johnson	Ergamisol	Levamisole	Cancer Therapy
Glaxo Wellcome	Eniluracil/776C85	5FU Enhancer	Cancer, Refractory Solid & Colorectal Cancer
Johnson & Johnson	Ergamisol	Levamisole	Colon Cancer
Rhone Poulenc	Campto	Irinotecan	Colorectal Cancer, Cervical Cancer
Pharmacia & Upjohn	Camptosar	Irinotecan	Colorectal Cancer, Cervical Cancer
Zeneca	Tomudex	Ralitrexed	Colorectal Cancer, Lung

- 113 -

Johnson & Johnson	Leustain	Cladribine	Cancer, Breast Cancer Hairy Cell Leukaemia
Ivax	Paxene	Paclitaxel	Kaposi Sarcoma
Sequus	Doxil	Doxorubicin, Liposomal	KS/Cancer
Sequus	Caelyx	Doxorubicin, Liposomal	KS/Cancer
Schering AG	Fludara	Fludarabine	Leukaemia
Pharmacia & Upjohn	Pharmorubicin	Epirubicin	Lung/Breast Cancer
Chiron	DepoCyt	DepoCyt	Neoplastic Meningitis
Zeneca	ZD1839	ZD 1839	Non Small Cell Lung Cancer, Pancreatic Cancer
BASF	LU 79553	Bis-Naphtalimide	Oncology
BASF	LU 103793	Dolastain	Oncology
Shering Plough	Caetyx	Doxorubicin-Liposome	Ovarian/Breast Cancer
Lilly	Gemzar	Gemcitabine	Pancreatic Cancer, Non Small Cell Lung Cancer, Breast, Bladder and Ovarian
Zeneca	ZD 0473/Anormed	ZD 0473/Anormed	Platinum based NSCL, ovarian etc.
Yamanouchi	YM 116	YM 116	Prostate Cancer
Nycomed Amersham	Seeds/I-125 Rapid St	Lodine Seeds	Prostate Cancer
Agouron	Cdk4/cdk2 inhibitors	cdk4/cdk2 inhibitors	Solid Tumors
Agouron	PARP inhibitors	PARP Inhibitors	Solid Tumors
Chiroscience	D4809	Dexifosamide	Solid Tumors
Bristol Myers Squib	UFT (Tegafur/Uracil)	UFT (Tegafur/Uracil)	Solid Tumors
Sankyo	Krestin	Krestin	Solid Tumors
Asta Medica	Ifex/Mesnex	Ifosamide	Solid Tumors
Bristol Meyers Squib	Ifex/Mesnex	Ifosamide	Solid Tumors
Bristol Myers Squib	Vumon	Teniposide	Solid Tumors
Bristol Myers Squib	Paraplatin	Carboplatin	Solid Tumors
Bristol Myers Squib	Plantinol	Cisplatin, Stealth	Solid Tumors
Bristol Myers Squib	Plantinol	Cisplatin	Solid Tumors
Bristol Myers Squib	Vepeside	Etoposide	Solid Tumors Melanoma
Zeneca	ZD 9331	ZD 9331	Solid Tumors, Advanced Colorectal
Chugai	Taxotere	Docetaxel	Solid Tumors, Breast Cancer
Rhone Poulenc	Taxotere	Docetaxel	Solid Tumors, Breast Cancer
Glaxo Wellcome	Prodrug of guanine arabinside	Prodrug of arabinside	T Cell Leukemia/Lymphoma & B Cell Neoplasm
Bristol Myers Squib	Taxane Analog	Taxane Analog	Taxol follow up

In one embodiment, the methods of the invention use immunostimulatory nucleic acids as a replacement to the use of IFN $\alpha$  therapy in the treatment of cancer. Currently, some treatment protocols call for the use of IFN $\alpha$ . Since IFN $\alpha$  is produced following the

- 114 -

administration of some immunostimulatory nucleic acids, these nucleic acids can be used to generate IFN $\alpha$  endogenously.

The invention also includes a method for inducing antigen non-specific innate immune activation and broad spectrum resistance to infectious challenge using the immunostimulatory nucleic acids. The term antigen non-specific innate immune activation as used herein refers to the activation of immune cells other than B cells and for instance can include the activation of NK cells, T cells or other immune cells that can respond in an antigen independent fashion or some combination of these cells. A broad spectrum resistance to infectious challenge is induced because the immune cells are in active form and are primed to respond to any invading compound or microorganism. The cells do not have to be specifically primed against a particular antigen. This is particularly useful in biowarfare, and the other circumstances described above such as travelers.

The stimulation index of a particular immunostimulatory nucleic acid can be tested in various immune cell assays. Preferably, the stimulation index of the immunostimulatory nucleic acid with regard to B cell proliferation is at least about 5, preferably at least about 10, more preferably at least about 15 and most preferably at least about 20 as determined by incorporation of  $^3\text{H}$  uridine in a murine B cell culture, which has been contacted with 20  $\mu\text{M}$  of nucleic acid for 20h at 37°C and has been pulsed with 1  $\mu\text{Ci}$  of  $^3\text{H}$  uridine; and harvested and counted 4h later as described in detail in PCT Published Patent Applications PCT/US95/01570 (WO 96/02555) and PCT/US97/19791 (WO 98/18810) claiming priority to U.S. Serial Nos. 08/386,063 and 08/960,774, filed on February 7, 1995 and October 30, 1997 respectively. For use *in vivo*, for example, it is important that the immunostimulatory nucleic acids be capable of effectively inducing an immune response, such as, for example, antibody production.

Immunostimulatory nucleic acids are effective in non-rodent vertebrate. Different immunostimulatory nucleic acid can cause optimal immune stimulation depending on the type of subject and the sequence of the immunostimulatory nucleic acid. Many vertebrates have been found according to the invention to be responsive to the same class of immunostimulatory nucleic acids, sometimes referred to as human specific immunostimulatory nucleic acids. Rodents, however, respond to different nucleic acids. As shown herein an immunostimulatory nucleic acid causing optimal



- 115 -

stimulation in humans may not generally cause optimal stimulation in a mouse and vice versa. An immunostimulatory nucleic acid causing optimal stimulation in humans often does, however, cause optimal stimulation in other animals such as cow, horses, sheep, etc. One of skill in the art can identify the optimal nucleic acid sequences useful for a particular species of interest using routine assays described herein and/or known in the art, using the guidance supplied herein.

The immunostimulatory nucleic acids may be directly administered to the subject or may be administered in conjunction with a nucleic acid delivery complex. A nucleic acid delivery complex shall mean a nucleic acid molecule associated with (e.g. ionically or covalently bound to; or encapsulated within) a targeting means (e.g. a molecule that results in higher affinity binding to target cell (e.g., B cell surfaces and/or increased cellular uptake by target cells). Examples of nucleic acid delivery complexes include nucleic acids associated with a sterol (e.g. cholesterol), a lipid (e.g. a cationic lipid, virosome or liposome), or a target cell specific binding agent (e.g. a ligand recognized by target cell specific receptor). Preferred complexes may be sufficiently stable *in vivo* to prevent significant uncoupling prior to internalization by the target cell. However, the complex can be cleavable under appropriate conditions within the cell so that the nucleic acid is released in a functional form.

Delivery vehicles or delivery devices for delivering antigen and nucleic acids to surfaces have been described. The Immunostimulatory nucleic acid and/or the antigen and/or other therapeutics may be administered alone (e.g., in saline or buffer) or using any delivery vehicles known in the art. For instance the following delivery vehicles have been described: Cochleates (Gould-Fogerite et al., 1994, 1996); Emulsomes (Vancott et al., 1998, Lowell et al., 1997); ISCOMs (Mowat et al., 1993, Carlsson et al., 1991, Hu et al., 1998, Morein et al., 1999); Liposomes (Childers et al., 1999, Michalek et al., 1989, 1992, de Haan 1995a, 1995b); Live bacterial vectors (e.g., *Salmonella*, *Escherichia coli*, *Bacillus calmatte-guerin*, *Shigella*, *Lactobacillus*) (Hone et al., 1996, Pouwels et al., 1998, Chatfield et al., 1993, Stover et al., 1991, Nugent et al., 1998); Live viral vectors (e.g., Vaccinia, adenovirus, Herpes Simplex) (Gallichan et al., 1993, 1995, Moss et al., 1996, Nugent et al., 1998, Flexner et al., 1988, Morrow et al., 1999); Microspheres (Gupta et al., 1998, Jones et al., 1996, Maloy et al., 1994, Moore et al., 1995, O'Hagan et al., 1994, Eldridge et al., 1989); Nucleic acid vaccines (Fynan et al., 1993, Kuklin et al.,

- 116 -

1997, Sasaki et al., 1998, Okada et al., 1997, Ishii et al., 1997); Polymers (e.g. carboxymethylcellulose, chitosan) (Hamajima et al., 1998, Jabbal-Gill et al., 1998); Polymer rings (Wyatt et al., 1998); Proteosomes (Vancott et al., 1998, Lowell et al., 1988, 1996, 1997); Sodium Fluoride (Hashi et al., 1998); Transgenic plants (Tacket et al., 1998, Mason et al., 1998, Haq et al., 1995); Virosomes (Gluck et al., 1992, Mengiardi et al., 1995, Cryz et al., 1998); Virus-like particles (Jiang et al., 1999, Leibl et al., 1998). Other delivery vehicles are known in the art and some additional examples are provided below in the discussion of vectors.

The term effective amount of a immunostimulatory nucleic acid refers to the amount necessary or sufficient to realize a desired biologic effect. For example, an effective amount of a immunostimulatory nucleic acid for inducing mucosal immunity is that amount necessary to cause the development of IgA in response to an antigen upon exposure to the antigen, whereas that amount required for inducing systemic immunity is that amount necessary to cause the development of IgG in response to an antigen upon exposure to the antigen. Combined with the teachings provided herein, by choosing among the various active compounds and weighing factors such as potency, relative bioavailability, patient body weight, severity of adverse side-effects and preferred mode of administration, an effective prophylactic or therapeutic treatment regimen can be planned which does not cause substantial toxicity and yet is entirely effective to treat the particular subject. The effective amount for any particular application can vary depending on such factors as the disease or condition being treated, the particular immunostimulatory nucleic acid being administered, the antigen, the size of the subject, or the severity of the disease or condition. One of ordinary skill in the art can empirically determine the effective amount of a particular immunostimulatory nucleic acid and/or antigen and/or other therapeutic agent without necessitating undue experimentation.

Subject doses of the compounds described herein for mucosal or local delivery typically range from about 0.1  $\mu$ g to 10 mg per administration, which depending on the application could be given daily, weekly, or monthly and any other amount of time therebetween. More typically mucosal or local doses range from about 10  $\mu$ g to 5 mg per administration, and most typically from about 100  $\mu$ g to 1 mg, with 2 - 4 administrations being spaced days or weeks apart. More typically, immune stimulant

- 117 -

doses range from 1  $\mu$ g to 10 mg per administration, and most typically 10 $\mu$ g to 1 mg, with daily or weekly administrations. Subject doses of the compounds described herein for parenteral delivery for the purpose of inducing an antigen-specific immune response, wherein the compounds are delivered with an antigen but not another therapeutic agent  
5 are typically 5 to 10,000 times higher than the effective mucosal dose for vaccine adjuvant or immune stimulant applications, and more typically 10 to 1,000 times higher, and most typically 20 to 100 times higher. Doses of the compounds described herein for parenteral delivery for the purpose of inducing an innate immune response or for increasing ADCC or for inducing an antigen specific immune response when the  
10 immunostimulatory nucleic acids are administered in combination with other therapeutic agents or in specialized delivery vehicles typically range from about 0.1  $\mu$ g to 10 mg per administration, which depending on the application could be given daily, weekly, or monthly and any other amount of time therebetween. More typically parenteral doses for these purposes range from about 10  $\mu$ g to 5 mg per administration, and most typically  
15 from about 100  $\mu$ g to 1 mg, with 2 - 4 administrations being spaced days or weeks apart. In some embodiments, however, parenteral doses for these purposes may be used in a range of 5 to 10,000 times higher than the typical doses described above.

For any compound described herein the therapeutically effective amount can be initially determined from animal models. A therapeutically effective dose can also be  
20 determined from human data for CpG oligonucleotides which have been tested in humans (human clinical trials have been initiated) and for compounds which are known to exhibit similar pharmacological activities, such as other mucosal adjuvants, e.g., LT and other antigens for vaccination purposes, for the mucosal or local administration. Higher doses are required for parenteral administration. The applied dose can be  
25 adjusted based on the relative bioavailability and potency of the administered compound. Adjusting the dose to achieve maximal efficacy based on the methods described above and other methods as are well-known in the art is well within the capabilities of the ordinarily skilled artisan.

The formulations of the invention are administered in pharmaceutically  
30 acceptable solutions, which may routinely contain pharmaceutically acceptable concentrations of salt, buffering agents, preservatives, compatible carriers, adjuvants, and optionally other therapeutic ingredients.

- 118 -

For use in therapy, an effective amount of the immunostimulatory nucleic acid can be administered to a subject by any mode that delivers the nucleic acid to the desired surface, e.g., mucosal, systemic. Administering the pharmaceutical composition of the present invention may be accomplished by any means known to the skilled artisan.

- 5 Preferred routes of administration include but are not limited to oral, parenteral, intramuscular, intranasal, intratracheal, inhalation, ocular, vaginal, and rectal.

- For oral administration, the compounds (i.e., immunostimulatory nucleic acids, antigens and other therapeutic agents) can be formulated readily by combining the active compound(s) with pharmaceutically acceptable carriers well known in the art. Such carriers enable the compounds of the invention to be formulated as tablets, pills, dragees, capsules, liquids, gels, syrups, slurries, suspensions and the like, for oral ingestion by a subject to be treated. Pharmaceutical preparations for oral use can be obtained as solid excipient, optionally grinding a resulting mixture, and processing the mixture of granules, after adding suitable auxiliaries, if desired, to obtain tablets or dragee cores.
- 10 Suitable excipients are, in particular, fillers such as sugars, including lactose, sucrose, mannitol, or sorbitol; cellulose preparations such as, for example, maize starch, wheat starch, rice starch, potato starch, gelatin, gum tragacanth, methyl cellulose, hydroxypropylmethyl-cellulose, sodium carboxymethylcellulose, and/or polyvinylpyrrolidone (PVP). If desired, disintegrating agents may be added, such as the cross-linked polyvinyl pyrrolidone, agar, or alginic acid or a salt thereof such as sodium alginate. Optionally the oral formulations may also be formulated in saline or buffers for neutralizing internal acid conditions or may be administered without any carriers.
- 15

- Dragee cores are provided with suitable coatings. For this purpose, concentrated sugar solutions may be used, which may optionally contain gum arabic, talc, polyvinyl pyrrolidone, carbopol gel, polyethylene glycol, and/or titanium dioxide, lacquer solutions, and suitable organic solvents or solvent mixtures. Dyestuffs or pigments may be added to the tablets or dragee coatings for identification or to characterize different combinations of active compound doses.
- 25

- Pharmaceutical preparations which can be used orally include push-fit capsules made of gelatin, as well as soft, sealed capsules made of gelatin and a plasticizer, such as glycerol or sorbitol. The push-fit capsules can contain the active ingredients in admixture with filler such as lactose, binders such as starches, and/or lubricants such as
- 30

- 119 -

talc or magnesium stearate and, optionally, stabilizers. In soft capsules, the active compounds may be dissolved or suspended in suitable liquids, such as fatty oils, liquid paraffin, or liquid polyethylene glycols. In addition, stabilizers may be added.

Microspheres formulated for oral administration may also be used. Such microspheres  
5 have been well defined in the art. All formulations for oral administration should be in dosages suitable for such administration.

For buccal administration, the compositions may take the form of tablets or lozenges formulated in conventional manner.

For administration by inhalation, the compounds for use according to the present  
10 invention may be conveniently delivered in the form of an aerosol spray presentation from pressurized packs or a nebulizer, with the use of a suitable propellant, *e.g.*, dichlorodifluoromethane, trichlorofluoromethane, dichlorotetrafluoroethane, carbon dioxide or other suitable gas. In the case of a pressurized aerosol the dosage unit may be determined by providing a valve to deliver a metered amount. Capsules and cartridges of  
15 *e.g.* gelatin for use in an inhaler or insufflator may be formulated containing a powder mix of the compound and a suitable powder base such as lactose or starch.

The compounds, when it is desirable to deliver them systemically, may be formulated for parenteral administration by injection, *e.g.*, by bolus injection or continuous infusion. Formulations for injection may be presented in unit dosage form,  
20 *e.g.*, in ampoules or in multi-dose containers, with an added preservative. The compositions may take such forms as suspensions, solutions or emulsions in oily or aqueous vehicles, and may contain formulatory agents such as suspending, stabilizing and/or dispersing agents.

Pharmaceutical formulations for parenteral administration include aqueous  
25 solutions of the active compounds in water-soluble form. Additionally, suspensions of the active compounds may be prepared as appropriate oily injection suspensions. Suitable lipophilic solvents or vehicles include fatty oils such as sesame oil, or synthetic fatty acid esters, such as ethyl oleate or triglycerides, or liposomes. Aqueous injection suspensions may contain substances which increase the viscosity of the suspension, such  
30 as sodium carboxymethyl cellulose, sorbitol, or dextran. Optionally, the suspension may also contain suitable stabilizers or agents which increase the solubility of the compounds to allow for the preparation of highly concentrated solutions.

Alternatively, the active compounds may be in powder form for constitution with a suitable vehicle, *e.g.*, sterile pyrogen-free water, before use.

The compounds may also be formulated in rectal or vaginal compositions such as suppositories or retention enemas, *e.g.*, containing conventional suppository bases such as cocoa butter or other glycerides.

In addition to the formulations described previously, the compounds may also be formulated as a depot preparation. Such long acting formulations may be formulated with suitable polymeric or hydrophobic materials (for example as an emulsion in an acceptable oil) or ion exchange resins, or as sparingly soluble derivatives, for example, as a sparingly soluble salt.

The pharmaceutical compositions also may comprise suitable solid or gel phase carriers or excipients. Examples of such carriers or excipients include but are not limited to calcium carbonate, calcium phosphate, various sugars, starches, cellulose derivatives, gelatin, and polymers such as polyethylene glycols.

Suitable liquid or solid pharmaceutical preparation forms are, for example, aqueous or saline solutions for inhalation, microencapsulated, encochleated, coated onto microscopic gold particles, contained in liposomes, nebulized, aerosols, pellets for implantation into the skin, or dried onto a sharp object to be scratched into the skin. The pharmaceutical compositions also include granules, powders, tablets, coated tablets, (micro)capsules, suppositories, syrups, emulsions, suspensions, creams, drops or preparations with protracted release of active compounds, in whose preparation excipients and additives and/or auxiliaries such as disintegrants, binders, coating agents, swelling agents, lubricants, flavorings, sweeteners or solubilizers are customarily used as described above. The pharmaceutical compositions are suitable for use in a variety of drug delivery systems. For a brief review of methods for drug delivery, see Langer, *Science* 249:1527-1533, 1990, which is incorporated herein by reference.

The immunostimulatory nucleic acids and optionally other therapeutics and/or antigens may be administered *per se* (neat) or in the form of a pharmaceutically acceptable salt. When used in medicine the salts should be pharmaceutically acceptable, but non-pharmaceutically acceptable salts may conveniently be used to prepare pharmaceutically acceptable salts thereof. Such salts include, but are not limited to, those prepared from the following acids: hydrochloric, hydrobromic, sulphuric, nitric,

phosphoric, maleic, acetic, salicylic, p-toluene sulphonic, tartaric, citric, methane sulphonic, formic, malonic, succinic, naphthalene-2-sulphonic, and benzene sulphonic. Also, such salts can be prepared as alkaline metal or alkaline earth salts, such as sodium, potassium or calcium salts of the carboxylic acid group.

5           Suitable buffering agents include: acetic acid and a salt (1-2% w/v); citric acid and a salt (1-3% w/v); boric acid and a salt (0.5-2.5% w/v); and phosphoric acid and a salt (0.8-2% w/v). Suitable preservatives include benzalkonium chloride (0.003-0.03% w/v); chlorobutanol (0.3-0.9% w/v); parabens (0.01-0.25% w/v) and thimerosal (0.004-0.02% w/v).

10           The pharmaceutical compositions of the invention contain an effective amount of a Immunostimulatory nucleic acid and optionally antigens and/or other therapeutic agents optionally included in a pharmaceutically-acceptable carrier. The term pharmaceutically-acceptable carrier means one or more compatible solid or liquid filler, diluents or encapsulating substances which are suitable for administration to a human or  
15 other vertebrate animal. The term carrier denotes an organic or inorganic ingredient, natural or synthetic, with which the active ingredient is combined to facilitate the application. The components of the pharmaceutical compositions also are capable of being commingled with the compounds of the present invention, and with each other, in a manner such that there is no interaction which would substantially impair the desired  
20 pharmaceutical efficiency.

The immunostimulatory nucleic acids useful in the invention may be delivered in mixtures with additional adjuvant(s), other therapeutics, or antigen(s). A mixture may consist of several adjuvants in addition to the Immunostimulatory nucleic acid or several antigens or other therapeutics.

25           A variety of administration routes are available. The particular mode selected will depend, of course, upon the particular adjuvants or antigen selected, the particular condition being treated and the dosage required for therapeutic efficacy. The methods of this invention, generally speaking, may be practiced using any mode of administration that is medically acceptable, meaning any mode that produces effective levels of an  
30 immune response without causing clinically unacceptable adverse effects. Preferred modes of administration are discussed above.

The compositions may conveniently be presented in unit dosage form and may be prepared by any of the methods well known in the art of pharmacy. All methods include the step of bringing the compounds into association with a carrier which constitutes one or more accessory ingredients. In general, the compositions are prepared by uniformly  
5 and intimately bringing the compounds into association with a liquid carrier, a finely divided solid carrier, or both, and then, if necessary, shaping the product. Liquid dose units are vials or ampoules. Solid dose units are tablets, capsules and suppositories. For treatment of a patient, depending on activity of the compound, manner of administration, purpose of the immunization (i.e., prophylactic or therapeutic), nature and severity of the  
10 disorder, age and body weight of the patient, different doses may be necessary. The administration of a given dose can be carried out both by single administration in the form of an individual dose unit or else several smaller dose units. Multiple administration of doses at specific intervals of weeks or months apart is usual for boosting the antigen-specific responses.

15 Other delivery systems can include time-release, delayed release or sustained release delivery systems. Such systems can avoid repeated administrations of the compounds, increasing convenience to the subject and the physician. Many types of release delivery systems are available and known to those of ordinary skill in the art. They include polymer base systems such as poly(lactide-glycolide), copolyoxalates,  
20 polycaprolactones, polyesteramides, polyorthoesters, polyhydroxybutyric acid, and polyanhydrides. Microcapsules of the foregoing polymers containing drugs are described in, for example, U.S. Patent 5,075,109. Delivery systems also include non-polymer systems that are: lipids including sterols such as cholesterol, cholesterol esters and fatty acids or neutral fats such as mono-di-and tri-glycerides; hydrogel release  
25 systems; sylastic systems; peptide based systems; wax coatings; compressed tablets using conventional binders and excipients; partially fused implants; and the like. Specific examples include, but are not limited to: (a) erosional systems in which an agent of the invention is contained in a form within a matrix such as those described in U.S. Patent Nos. 4,452,775, 4,675,189, and 5,736,152, and (b) diffusional systems in which  
30 an active component permeates at a controlled rate from a polymer such as described in U.S. Patent Nos. 3,854,480, 5,133,974 and 5,407,686. In addition, pump-based hardware delivery systems can be used, some of which are adapted for implantation.



- 123 -

The present invention is further illustrated by the following Examples, which in no way should be construed as further limiting. The entire contents of all of the references (including literature references, issued patents, published patent applications, and co-pending patent applications) cited throughout this application are hereby  
5 expressly incorporated by reference.

### Examples

#### ***Materials and Methods:***

**Oligodeoxynucleotides:** Native phosphodiester and phosphorothioate-modified ODN were purchased from Operon Technologies (Alameda, CA) and Hybridon  
10 Specialty Products (Milford, MA). ODN were tested for endotoxin using the LAL-assay (LAL-assay BioWhittaker, Walkersville, MD; lower detection limit 0.1 EU/ml). For *in vitro* assays, ODN were diluted in TE-buffer (10 mM Tris, pH 7.0, 1 mM EDTA), and stored at -20° C. For *in vivo* use, ODN were diluted in phosphate buffered saline (0.1 M PBS, pH 7.3) and stored at 4°C. All dilutions were carried out using pyrogen-free  
15 reagents.

**Isolation of human PBMC and cell culture:** Peripheral blood mononuclear cells (PBMC) were isolated from peripheral blood of healthy volunteers by Ficoll-Paque density gradient centrifugation (Histopaque-1077, Sigma Chemical Co., St. Louis, MO) as described (Hartmann et al., 1999 Proc. Natl. Acad. Sci USA 96:9305-10). Cells were  
20 suspended in RPMI 1640 culture medium supplemented with 10% (v/v) heat-inactivated (56°C, 1 h) FCS (HyClone, Logan, UT), 1.5 mM L-glutamine, 100 U/ml penicillin and 100 µg/ml streptomycin (all from Gibco BRL, Grand Island, NY) (complete medium). Cells (final concentration  $1 \times 10^6$  cells/ml) were cultured in complete medium in a 5% CO<sub>2</sub> humidified incubator at 37°C. ODN and LPS (from *Salmonella typhimurium*, Sigma  
25 Chemical Co., St. Louis, MO) or anti-IgM were used as stimuli. For measurement of human NK lytic activity, PBMC were incubated at  $5 \times 10^6$ /well in 24-well plates. Cultures were harvested after 24 hours, and cells were used as effectors in a standard 4 hours <sup>51</sup>Cr-release assay against K562 target cells as previously described (Ballas et al., 1996 J. Immunol. 157:1840-1845). For B cell proliferation, 1 µCi of <sup>3</sup>H thymidine was  
30 added 18 hours before harvest, and the amount of <sup>3</sup>H thymidine incorporation was determined by scintillation counting at day 5. Standard deviations of the triplicate wells were < 5%.

- 124 -

**Flow cytometry on human PBMC:** Surface antigens on primate PBMC were stained as previously described (Hartmann et al., 1998 J. Pharmacol. Exp. Ther. 285:920-928). Monoclonal antibodies to CD3 (UCHT1), CD14 (M5E2), CD19 (B43), CD56 (B159), CD69 (FN50) and CD86 (2331 [FUN-1]) were purchased from Pharmingen, San Diego, CA. IgG<sub>1</sub>, $\kappa$  (MOPC-21) and IgG<sub>2b</sub>, $\kappa$  (Hartmann et al., 1999 Proc. Natl. Acad. Sci USA 96:9305-10) were used to control for non-specific staining. NK cells were identified by CD56 expression on CD3, CD14 and CD19 negative cells, whereas B cells were identified by expression of CD19. Flow cytometric data of 10000 cells per sample were acquired on a FACScan (Beckton Dickinson Immunocytometry Systems, San Jose, CA). The viability of cells within the FSC/SSC gate used for analysis was examined by propidium iodide staining (2  $\mu$ g/ml) and found to be higher than 98%. Data were analyzed using the computer program FlowJo (version 2.5.1, Tree Star, Inc., Stanford, CA).

**Results:**

*Example 1: CpG-dependent stimulation of human B cells depends on methylation and ODN length.*

Human PBMC were obtained from normal donors and cultured for five days at  $2 \times 10^5$  cells/well with the indicated concentrations of the indicated ODN sequences. As shown in Table F, human PBMCs proliferate above the background when cultured with a variety of different CpG ODN, but also show some proliferation even with ODN that do not contain any CpG motifs. The importance of unmethylated CpG motifs in providing optimal immune stimulation with these ODN is demonstrated by the fact that ODN 1840 (SEQ ID NO. 83) induces 56,603 counts of <sup>3</sup>H-thymidine incorporation whereas the same T-rich ODN with the CpG motifs methylated (non-CpG), 1979 (SEQ ID NO. 222), induces lower, but still increased over background, activity (only 18,618 counts) at the same concentration of 0.6  $\mu$ g/ml. The reduced proliferation at higher ODN concentrations may be an artifact of the cells becoming exhausted under these experimental conditions or could reflect some toxicity of the higher ODN concentrations. Interestingly, shorter ODN containing CpG motifs, such as the 13-14 mers 2015 and 2016, are less stimulatory despite the fact that their molar concentration would actually be higher since the ODNs were added on the basis of mass rather than molarity. This demonstrates that ODN length may also be an important determinant in the immune

- 125 -

effects of the ODN. A non-CpG ODN but slight T-rich ODN (about 30% T), 1982 (SEQ ID NO. 225), caused only a small amount of background cell proliferation.

**Table F**

<b>Oligo Concentration</b>			
<b>ODN#</b>	<b>0.15 µg/ml</b>	<b>0.6 µg/ml</b>	<b>2 µg/ml</b>
Cues only	648	837	799
1840 (SEQ ID NO. 83)	5744	56,603	31,787
2016 (SEQ ID NO. 256)	768	4607	20,497
1979 (SEQ ID NO. 222)	971	18,618	29,246
1892 (SEQ ID NO. 135)	787	10,078	22,850
2010 (SEQ ID NO. 250)	849	20,741	8,054
2012 (SEQ ID NO. 252)	2586	62,955	52,462
2013 (SEQ ID NO. 253)	1043	47,960	47,231
2014 (SEQ ID NO. 254)	2700	50,708	46,625
2015 (SEQ ID NO. 255)	1059	23,239	36,119

Numbers represent cpm of <sup>3</sup>H-thymidine incorporation for cultures of human

5 PBMCs set up as described above.

*Example 2. Concentration-dependent activation of human NK cell activity with thymidine-rich ODN.*

- 126 -

Human PBMCs were cultured for 24 hours with a panel of different CpG or non-CpG ODN at two different concentrations, and then tested for their ability to kill NK target cells as described previously (Ballas et al., 1996 J. Immunol. 157:1840-1845). Killing is measured as lytic units, or L.U. The human donor used in this experiment had a background level of 3.69 L.U. which increased to 180.36 L.U. using the positive control, IL-2. A CpG oligo, 2006 (SEQ ID NO. 246), induced high levels of NK lytic function at a low concentration of 0.6, and a lower level at a concentration of 6.0. Surprisingly, a T-rich ODN in which the CpG motifs of 2006 were methylated (ODN at 2117 (SEQ ID NO. 358)) or inverted to GpCs (ODN 2137 (SEQ ID NO. 886)) retained strong immune stimulatory function at the higher ODN concentrations, as shown in Table G. These concentration-dependent immune stimulatory effects are not a general property of the phosphorothioate backbone since the experiments described below demonstrate that a poly-A ODN, is nonstimulatory above background levels. Some stimulation is seen with a 24-base long ODN in which all of the base positions are randomized so that A, C, G, and T will occur at a frequency of 25% in each of the base positions (ODN 2182 (SEQ ID NO. 432)). However, the stimulatory effect of such a 24-base ODN is greatly enhanced if it is pure poly-T, in which case stimulation is also seen at the lowest concentration of 0.6 µg/ml (ODN 2183 (SEQ ID NO. 433)). In fact, the stimulatory activity of ODN SEQ ID NO. 433 at this low concentration is higher than that of any other ODN tested at this low concentration, aside from the optimal human immune stimulatory ODN of SEQ ID NO. 246. In fact, the higher concentration of ODN SEQ ID NO. 433 stimulated more NK activity than any other phosphorothioate ODN except for the strong CpG ODN 2142 (SEQ ID NO. 890), which was marginally higher. If the G content of ODN SEQ ID NO. 246 is increased relative to the T content by addition of more Gs, thus resulting in a decrease in the proportion of T nucleotides the immune stimulatory effect of the ODN is reduced (see ODN 2132 (SEQ ID NO. 373)). Thus, the T content of an ODN is an important determinant of its immune stimulatory effect. Although a poly-T ODN is the most stimulatory of the non-CpG ODN, other bases are also important in determining the immune stimulatory effect of a non-CpG ODN. ODN 2131 (SEQ ID NO. 372), in which slightly more than half of the bases are T and in which there are no Gs, is immune stimulatory at a concentration of 6 µg/ml but has less activity than other T-rich ODN. If the 6 A's in ODN 2131 (SEQ ID NO. 372)

- 127 -

are replaced by 6 Gs, the immune stimulatory effect of the ODN can be increased (see ODN 2130 (SEQ ID NO. 371)).

**Table G*****HUMAN PBL CULTURED OVERNIGHT WITH OLIGOS***

MR	3605						
SR	256						
%SR	7.11						
EFFECTOR	0.63	1.25	2.50	5.00	10.00	20.00	
<b>CONTROL [RM]</b>							<b>L.U.</b>
ALONE	2.65	5.45	10.15	17.65	29.92	39.98	<b>3.69</b>
+ IL2 (100 U/ml)	35.95	57.66	86.26	100.39	99.71	93.64	<b>180.36</b>
+ 1585 (0.6 ug/ml)	3.75	6.10	12.14	23.70	36.06	43.98	<b>5.48</b>
+ 1585 (6.0 ug/ml)	15.42	31.09	47.07	73.34	94.29	97.73	<b>35.85</b>
+ 2006 (0.6 ug/ml)	6.71	15.99	26.92	44.75	64.12	68.83	<b>16.96</b>
+ 2006 (6.0 ug/ml)	6.19	8.18	16.13	24.35	39.35	56.07	<b>8.04</b>
+ 2117 (0.6 ug/ml)	4.54	4.73	9.56	18.04	28.57	39.85	<b>3.49</b>
+ 2117 (6.0 ug/ml)	7.03	10.76	16.90	30.59	52.14	59.46	<b>10.96</b>
+ 2137 (0.6 ug/ml)	4.61	5.35	10.04	15.16	23.79	37.86	<b>2.57</b>
+ 2137 (6.0 ug/ml)	7.99	10.37	16.55	32.32	49.78	60.30	<b>11.01</b>
+ 2178 (0.6 ug/ml)	2.88	4.52	11.47	16.05	24.85	34.27	<b>2.37</b>
+ 2178 (6.0 ug/ml)	4.21	5.03	11.16	16.39	28.22	36.45	<b>2.94</b>
+ 2182 (0.6 ug/ml)	2.42	6.57	10.49	19.73	26.55	35.30	<b>2.89</b>
+ 2182 (6.0 ug/ml)	4.11	7.98	14.60	26.56	40.40	51.98	<b>7.59</b>

- 128 -

+ 2183 (0.6 ug/ml)	3.73	8.46	15.52	24.48	37.78	56.77	7.80
+ 2183 (0.6 ug/ml)	8.86	12.89	23.08	41.49	66.26	75.85	16.57
+ 2140 (0.6 ug/ml)	3.78	5.27	12.30	20.79	35.75	45.62	5.40
+ 2140 (6.0 ug/ml)	6.56	13.24	21.26	37.96	60.80	73.05	14.82
+ 2141 (0.6 ug/ml)	2.63	6.34	10.21	17.73	30.93	43.57	4.29
+ 2141 (6.0 ug/ml)	4.98	15.30	25.22	37.88	58.47	69.12	14.83
+ 2142 (0.6 ug/ml)	3.18	3.66	6.99	14.62	19.68	32.52	1.56
+ 2142 (6.0 ug/ml)	7.08	15.80	25.65	41.72	68.09	73.14	17.11
+ 2143 (0.6 ug/ml)	4.12	6.90	10.77	22.96	35.78	42.94	5.19
+ 2143 (6.0 ug/ml)	3.16	8.40	12.38	21.69	34.80	54.21	6.64
+ 2159 (6.0 ug/ml)	5.05	11.76	21.67	41.12	51.68	65.47	13.19
+ 2132 (6.0 ug/ml)	4.23	6.06	10.50	18.74	32.68	44.06	4.61
+ 2179 (6.0 ug/ml)	6.14	9.49	21.06	42.48	60.12	71.87	14.54
+ 2180 (6.0 ug/ml)	2.37	8.57	15.44	29.66	44.35	61.31	9.47
+ 2133 (6.0 ug/ml)	6.53	12.58	23.10	38.03	61.16	68.36	14.62
+ 2134 (6.0 ug/ml)	7.51	12.14	21.14	32.46	54.47	67.12	12.98
+ 2184 (6.0 ug/ml)	5.22	9.19	17.54	30.76	45.35	63.55	10.42
+ 2185 (6.0 ug/ml)	8.11	14.77	26.27	40.31	55.61	70.65	15.60
+ 2116 (6.0 ug/ml)	5.58	10.54	16.77	37.82	59.80	66.33	13.07
+ 2181 (6.0 ug/ml)	4.43	9.85	17.55	27.05	53.16	69.16	11.43
+ 2130 (6.0 ug/ml)	3.81	8.07	17.11	27.17	42.04	53.73	8.27

- 129 -

+ 2131 (6.0 ug/ml)	2.29	6.73	7.30	18.02	32.73	49.06	<b>5.08</b>
+ 2156 (0.3 ug/ml)	2.50	5.26	8.20	15.95	26.64	33.07	<b>2.31</b>
+ 2156 (1.0 ug/ml)	5.91	10.99	17.31	26.97	50.64	63.78	<b>10.84</b>
+ 2157 (0.3 ug/ml)	2.36	4.00	6.65	12.94	24.13	38.86	<b>2.58</b>
+ 2157 (1.0 ug/ml)	3.72	9.55	17.15	34.55	52.27	65.33	<b>11.58</b>
+ 2158 (0.3 ug/ml)	1.25	2.36	6.90	16.39	15.63	29.82	<b>1.17</b>
+ 2158 (1.0 ug/ml)	4.73	7.26	11.07	15.55	30.80	43.71	<b>4.16</b>
+ 2118 (0.6 ug/ml)	1.55	3.38	6.85	13.36	20.15	27.71	<b>1.13</b>
+ 2118 (6.0 ug/ml)	2.65	3.88	9.29	12.19	22.47	28.99	<b>1.34</b>

*Example 3: Induction of B cell proliferation by T-rich non-CpG ODN.*

To assess the ability of T-rich ODN to activate B cell proliferation, human PBMCs were stained with the cytoplasmic dye CFSE, incubated with five days with the indicated ODN at either 0.15 or 0.3 ug/ml, and then analyzed by flow cytometry. B cells were identified by gating on cells positive for the lineage marker CD19). CpG ODN 2006 was a strong inducer of B cell proliferation, and this effect was reduced if the CpG motifs were methylated or inverted to GpC as shown in Figure 1 at an ODN concentration of 0.3 ug/ml. The base composition of the ODN appears to be important in determining the immune stimulatory effect. Reducing the T content of an ODN substantially reduces immune stimulatory effect, as exemplified by ODN 2177 (SEQ ID NO. 427) in which 6 of the Ts present in ODN 2137 (SEQ ID NO. 886) have been switched to A's, resulting in a greatly reduced immune stimulatory effect. The importance of T's in the immune stimulatory effect of an ODN is also shown by comparison of ODN 2116 (SEQ ID NO. 357) and 2181 (SEQ ID NO. 431), which differ in the 3' end of the ODN. ODN 2181, in which the 3' end is poly-T is more stimulatory

- 130 -

than ODN 2116, in which the 3' end is poly-C, despite the fact that both ODN have a TCGTCG at the 5' end.

*Example 4: B Cell Proliferation Induced by TG Oligonucleotides*

The stimulatory effects of TG motifs are shown in Figure 2. ODN 2137 has the  
5 identical base composition as ODN 2006, but the CG motifs have all been inverted to  
GC's resulting in a CG-free nucleic acid. ODN does however contain 6 TG  
dinucleotides. In ODN 2177, all the TG dinucleotides of ODN 2137 have been changed  
to AG. Although ODN 2177 contains only 6 adenines, it is virtually nonstimulatory at a  
concentration of 0.2 µg/ml. For comparison, an ODN 24 bases in length in which each  
10 position is randomized to be any of the four bases (ODN 2182) induces > 12% of B cells  
to proliferate at a concentration of 0.2 µg/ml. These results indicate that the stimulatory  
effects of ODN 2137 are not simply those of a phosphorothioate backbone, but relate to  
the presence of TG dinucleotides.

In order to determine the effect of varying the number of TG dinucleotide motifs,  
15 ODN 2200 and ODN 2202 were compared, as shown in Figure 2. Both ODN contain 18  
Ts and 6 Gs, but in ODN 2200 all of the Gs are consecutive, so that there is only one TG  
dinucleotide, whereas in ODN 2202, the Gs are split up as GG dinucleotides throughout  
the ODN so that there are three TGs. ODN 2202 is significantly more stimulatory than  
ODN 2200, consistent with the model that at least three TG motifs in an ODN are  
20 required for optimal stimulatory activity. It is likely that even higher levels of  
stimulation could be achieved if the TG motifs had been optimized as taught herein.

*Example 5: Effects of TTG versus TTG motifs.*

Figure 3 shows the results of experiments conducted to study TG content in terms  
of the relative levels of Ts versus Gs as it relates to the stimulatory effect of an ODN.  
25 The Figure shows that an ODN in which all of the bases are randomized to be either T or  
G (ODN 2188 (SEQ ID NO. 905)) is nonstimulatory at a concentration of 0.2 µg/ml,  
similar to an ODN in which all of the bases are randomized to be either A or G (ODN  
2189 (SEQ ID NO. 906)). However, at the higher concentration of 2 µg/ml, the  
randomized T/G ODN 2188 is significantly more stimulatory. This latter level of  
30 stimulation is still lower than that which occurs with a totally randomized ODN (ODN  
2182 (SEQ ID NO. 432)). The highest stimulation at low concentrations is seen with an



ODN in which half of the bases are fixed at T and the other half of the bases are randomized to be either T or G (ODN 2190 (SEQ ID NO. 907)). Since every other base is fixed to be a T, there cannot be any TG motifs. The data in Figure 3 show that increasing the TG content of an ODN improves its stimulatory activity.

5 In yet other experiments, the results of which are not diagrammed herein, ODN 2190 (SEQ ID NO. 907) exhibited a stimulation of NK activity compared to ODN 2188 (SEQ ID NO. 905) or ODN 2189 (SEQ ID NO. 906).

### Examples 6-8

**Introduction:**

10       Above, we demonstrated that Poly T sequences are able to enhance stimulation of B and NK cells. Here and below we investigate the effect of a variety of non-CpG T-rich ODN as well as Poly C ODN for their ability to stimulate human B cells, NK cells and monocytes.

### Materials and Methods:

[illegible]

**Cell preparation and cell culture:** Human PBMC were isolated from peripheral blood of healthy volunteers, obtained by the German Red Cross (Ratingen, Germany), as described above in Example 1, but all material were purchased from Life Technologies, Germany and were endotoxin-tested. For the B cell, NK cell and monocyte activation assays PBMC were cultured in complete medium at a concentration of  $2 \times 10^6$  cells/ml in 200 $\mu$ l in 96 round bottom plates in a humidified incubator at 37°C. Different ODNs, LPS (Sigma) or IL-2 (R&D Systems, USA) were used as stimuli. At the indicated time points, cells were harvested for flow cytometry.

**Flow cytometry:** MAbs used for staining of surface antigens were: CD3, CD14, CD19, CD56, CD69, CD80 and CD86 (all obtained from Pharmingen/Becton Dickinson, Germany). For monocytes Fc receptors were blocked using human IgG (Myltenyi, Germany) as previously described (Bauer, M et al 1999 *Immunology* 97:699). Flow cytometric data of at least 1000 cells of a specified subpopulation (B cells, monocytes, NK cells, NKT cells or T cells) were acquired on a FACSCalibur (Becton Dickinson). Data were analyzed using the program CellQuest (Becton Dickinson).

**NK-mediated cytotoxicity:** PBMC were cultured overnight with or without 6 $\mu$ g/ml ODN or 100U/ml IL-2 at 37°C, 5% CO<sub>2</sub>. The next morning, K-562 target cells were labeled with a fluorescent dye, CFSE, as described previously for human B cells (Hartmann, G., and A. M. Krieg. 2000 *J. Immunol.* 164:944). PBMC were added in different ratios (50:1, 25:1 and 12.5:1) to  $2 \times 10^5$  target cells and incubated for 4h at 37°C. Cells were harvested and incubated with the DNA-specific dye 7-AAD (Pharmingen) for detection of apoptotic cells. Results were measured by flow cytometry.

**ELISA:** PBMC ( $3 \times 10^6$  cells/ml) were cultured with the specified concentrations of ODN or LPS for 24h (IL-6, IFN $\gamma$  and TNF $\alpha$ ) or 8h (IL-1 $\beta$ ) in 48 well plates in a humidified atmosphere at 37°C. Supernatants were collected and cytokines were measured using OPTeia ELISA Kits (Pharmingen) for IL-6, IFN $\gamma$  and TNF $\alpha$  or an Eli-pair ELISA assay (Hoelzel, Germany) for IL-1 $\beta$  according to the manufacturer protocols.

***Example 6: B cell activation induced by ODNs lacking CpG motifs***

In the Experiments described above in Example 3, we demonstrate that T-rich ODN were capable of activating B cells. We expand those studies here using additional ODN and different cell and reagent sources. In a first set of experiments, we compared the activation potential of different non-CpG T-rich ODNs with the very potent known

- 133 -

CpG ODN 2006 (SEQ ID NO.: 246). PBMC ( $2 \times 10^6$  cells/ml) of a blood donor ( $n=2$ ) were incubated with the indicated concentrations of ODNs 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ ID NO.: 886), 5126 (SEQ ID NO.: 1058), and 5162 (SEQ ID NO.: 1094). Cells were incubated for 48h at 37°C as described above and  
5 stained with mAb for CD19 (B cell marker) and CD86 (B cell activation marker, B7-2). Expression was measured by flow cytometry.

Using different concentrations of ODNs, we showed (Fig. 4) that T-rich ODNs without a CpG motif, can induce stimulation of human B cells. ODN 5126 (SEQ ID NO.: 1058) which contains only a single poly-T sequence but is greater than 50% T,  
10 caused high levels of human B cell activation. Although there are some similarities to SEQ ID NO.: 246 (e.g. more than 80% T/G content), this ODN clearly lacks any known immunostimulatory CpG motif. Surprisingly, for all tested T-rich ODNs, the highest stimulatory index was obtained at concentrations between 3 and 10 µg/ml. The highest stimulatory index of the tested ODNs was achieved by CpG/T-rich ODN SEQ ID NO.:  
15 246 at 0.4 µg/ml. Interestingly, the activity decreased at high concentrations.

Poly A, Poly C and Poly T sequences were synthesized and tested for biological activity. PBMC ( $2 \times 10^6$  cells/ml) of one representative donor ( $n=3$ ) were stimulated as described above by 0.4 µg/ml, 1.0 µg/ml or 10.0 µg/ml of the following ODNs: 2006 (SEQ ID NO.: 246), 2196 (SEQ ID NO.: 913) (Poly T, 18 bases), 2194 (SEQ ID NO.: 911)  
20 (Poly T, 27 bases), 5162 (SEQ ID NO.: 1094) (Poly T, 30 bases), 5163 (SEQ ID NO.: 1095) (Poly A, 30 bases), 5168 (SEQ ID NO.: 1096) (Poly C, 30 bases) and 5169 (SEQ ID NO.: 1097) (Poly CG, 30 bases). Expression of the activation marker CD86 (B7-2) on CD19-positive B cells was measured by flow cytometry.

Fig. 5 demonstrates that the length of the sequence, at least for Poly T ODNs, has  
25 an important impact on its activity. A Poly T sequence containing only 18 bases (SEQ ID NO.: 913) was shown to be less stimulatory than one with 27 bases (SEQ ID NO.: 911) or one with 30 bases (SEQ ID NO.: 1094) with a clear rank of stimulation: SEQ ID NO.: 1094 > SEQ ID NO.: 911 > SEQ ID NO.: 913. Poly A (SEQ ID NO.: 1095) or Poly CG (SEQ ID NO.: 1097) sequences, in contrast, do not induce activation of human B  
30 cells. Surprisingly it was also discovered that Poly C sequences (SEQ ID NO.: 1096) can activate human B cells at least at high concentrations (10 µg/ml) (Fig. 5).

Two other T-rich ODNs, namely 1982 (SEQ ID NO.: 225) and 2041 (SEQ ID NO.: 282) lacking CpG motifs were tested for their effect on human B cells. PBMC (n=2) were incubated with the indicated concentrations of ODN 2006 (SEQ ID NO.: 246), 1982 (SEQ ID NO.: 225) and 2041 (SEQ ID NO.: 282) as described above. B cell activation (expression of the activation marker CD86) was measured by flow cytometry.

Fig. 6 demonstrates that T-rich non-CpG ODN are immunostimulatory at concentrations higher than 1 µg/ml. Incorporation of a CpG motif into 1982 enhanced the immunostimulatory activity. Elongation with a Poly T sequence did not enhance the immunostimulatory activity of this already T-rich ODN but rather, decreased the activation potential slightly.

***Example 7: Immunostimulation of non-CpG ODNs is reflected in the enhancement of NK activation, NK cytotoxicity and monocyte activation***

NK cells as well as monocytes were tested for their response to non-CpG ODNs. PBMC ( $2 \times 10^6$  cells/ml) were incubated with 6 µg/ml of the following ODNs (n=4): 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ ID NO.: 886), 2183 (SEQ ID NO.: 433), 2194 (SEQ ID NO.: 911) and 5126 (SEQ ID NO.: 1058). After 24h of cultivation at 37°C cells were harvested and stained with mAb for CD3 (T cell marker), CD56 (NK cell marker) and CD69 (early activation marker) as described above. Expression of CD69 on CD56-positive NK cells was measured by flow cytometry.

Fig. 7 shows that for Poly T ODNs similar effects can be observed as described in Fig. 5. The stimulation of NK cells, like B cells, may be influenced by the length of the ODN. ODN 2183 (SEQ ID NO.: 433) (21 bases) induced activation of NK cells but to a lesser extent than the longer ODN 2194 (SEQ ID NO.: 911) (27 bases), as measured by enhanced expression of the early activation marker CD69. ODN 5126 (SEQ ID NO.: 1058) was also demonstrated to activate human NK cells (Fig. 7).

It is believed that the anti-tumor activity of CpG ODNs can be assessed by the ability of the ODN to enhance NK-mediated cytotoxicity *in vitro*. ODNs containing at the 5' and 3' ends stretches of Poly G were shown to result in the highest induction of cytotoxicity (Ballas, Z. K., et al. 1996 *J. Immunol.* 157:1840). To investigate the influence of non-CpG T-rich ODN on NK cytotoxicity, we analyzed the effect of the

- 135 -

ODNs 2194 (SEQ ID NO.: 911) and 5126 (SEQ ID NO.: 1058) on NK-mediated lysis (Fig. 8). NK-mediated lysis of K-562 target cells was measured after over night incubation of PBMC with 6µg/ml of the ODN 2006 (SEQ ID NO.: 246), SEQ ID NO.: 911 (SEQ ID NO.: 911 ) (Poly T, 27 bases) and 5126 (SEQ ID NO.: 1058) as described  
5 above. SEQ ID NO.: 1058 demonstrated small increases in lysis by human NK cells as compared to no ODN. SEQ ID NO.: 911 and SEQ ID NO.: 246 enhanced human NK cell cytotoxicity to an even higher extent.

Previous reports demonstrated that not only NK cells but also NKT cells are mediators of cytotoxic responses to tumor cells (14). We, therefore, looked at the  
10 potential activation of human NKT cells by T-rich non-CpG ODN. PBMC of one representative donor (n=2) were incubated with 6µg/ml ODN 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ ID NO.: 886), 2183 (SEQ ID NO.: 433), 2194 (SEQ ID NO.: 913) and 5126 (SEQ ID NO.: 1058) for 24h as described above. Activation of NKT cells was measured by flow cytometry after staining of cells with  
15 mAb for CD3 (T cell marker), CD56 (NK cell marker) and CD69 (early activation marker). Shown is the expression of CD69 on CD3 and CD56 double-positive cells (NKT cells).

In Fig. 9, SEQ ID NO.: 911 as well as SEQ ID NO.: 1058 were found to stimulate NKT cells. Similar to NK cells SEQ ID NO.: 911 (Poly T) was more active  
20 than SEQ ID NO. 1058. In addition, as described above for B cells and NK cells, the length of the ODN has some influence on the immunostimulatory potential, with the longer ODN having stronger effects on NKT cells. Similar results were observed for human T cells.

Another type of cell of the immune system involved in fighting infections is the  
25 monocytes. These cells release upon activation a variety of cytokines and can mature into dendritic cells (DC), professional antigen-presenting cells (Roitt, I., J. Brostoff, and D. Male. 1998. *Immunology*. Mosby, London). Fig. 10 shows activation of human monocytes after culturing of PBMC with different ODNs. PBMC ( $2 \times 10^6$  cells/ml) were incubated with 6µg/ml 2006 (SEQ ID NO.: 246), 2117 (SEQ ID NO.: 358), 2137 (SEQ  
30 ID NO.: 886), 2178 (SEQ ID NO.: 1096), 2183 (SEQ ID NO.: 433), 2194 (SEQ ID NO.: 911), 5126 (SEQ ID NO.: 1058) and 5163 (SEQ ID NO.: 1095) overnight at 37°C as

- 136 -

described above (n=3). Cells were harvested and stained for CD14 (monocyte marker) and CD80 (B7-1, activation marker). Expression was measured by flow cytometry.

As demonstrated above for NK and B cells, T-rich sequences (e.g., SEQ ID NO.: 433, SEQ ID NO.: 911) of different length induce monocyte stimulation but have  
5 different levels of activity e.g., SEQ ID NO.: 433 > SEQ ID NO.: 911. Poly A (SEQ ID NO.: 1095) as well as Poly C (SEQ ID NO.: 1096 (2178) sequences, in contrast, did not lead to activation of monocytes (measured by the upregulation of CD80 at a concentration of 6µg/ml ODN).

***Example 8: Induction of cytokine release by non-CpG ODNs***

10 Next the ability of different T-rich ODNs to influence the cytokine milieu was examined. PBMC ( $3 \times 10^6$  cells/ml) were cultured for 24h with or without 6µg/ml of the indicated ODNs or 1µg/ml LPS as positive control (n=2). After incubation supernatants were collected and TNFα measured by ELISA as described above and the results are shown in Fig. 11. PBMC were cultured with the indicated ODNs (1.0µg/ml) as  
15 described in Fig. 11 and IL-6 was measured in the supernatants by ELISA and the results are shown in Fig. 12.

Fig. 11 and 12 demonstrate that T-rich non-CpG and T-rich/CpG ODNs can induce the secretion of the pro-inflammatory cytokines TNFα and IL-6. For both cytokines, ODN 5126 (SEQ ID NO.: 1058) was found in most assays to be as potent as  
20 ODN 2194 (SEQ ID NO.: 911). It is known that CpG ODNs influence the Th1/Th2 balance by preferentially inducing Th1 cytokines (Krieg, A. M. 1999 *Biochemica et Biophysica Acta* 93321:1). To test whether T-rich ODN caused a similar shift to Th1 cytokines, IFNγ production in PBMC was measured. In a first set of experiments, it was demonstrated that, as described for IL-6 and TNFα, ODNs SEQ ID NO.: 1058 and SEQ  
25 ID NO.: 911 induced the release of comparable amounts of this Th1 cytokine IFNγ. In addition, it was demonstrated that another pro-inflammatory cytokine, IL-1β, was released upon culture of PBMC with these two ODNs. Although the amount of these cytokines induced by the T-rich ODN lacking CpG motifs was less than when CpG ODN SEQ ID NO.: 246 was used the amounts induced by T-rich ODN were significantly  
30 higher than the control.

**Examples 9-11*****Introduction:***

An optimal CpG motif for immune system activation in non-rodent vertebrates is described herein. A phosphodiester oligonucleotide containing this motif was found to strongly stimulate CD86, CD40, CD54 and MHC II expression, IL-6 synthesis and proliferation of primary human B-cells. These effects required internalisation of the oligonucleotide and endosomal maturation. This CpG motif was associated with the sustained induction of the NF $\kappa$ B p50/p65 heterodimer and of the transcription factor complex activating protein-1 (AP-1). Transcription factor activation by CpG DNA was preceded by increased phosphorylation of the stress kinases c-jun NH<sub>2</sub> terminal kinase (JNK) and p38, and of activating transcription factor-2 (ATF-2). In contrast to CpG, signaling through the B-cell receptor led to activation of extracellular receptor kinase (ERK) and to phosphorylation of a different isoform of JNK.

***Materials and Methods:***

**Oligodeoxynucleotides:** Unmodified (phosphodiester, PE) and modified nuclease-resistant (phosphorothioate, PS) ODN were purchased from Operon Technologies (Alameda, CA) and Hybridon Specialty Products (Milford, MA). The sequences used are provided in Table H. *E. coli* DNA and calf thymus DNA were purchased from Sigma Chemical Co., St. Louis, MO. Genomic DNA samples were purified by extraction with phenol-chloroform-isoamyl alcohol (25/24/1) and ethanol precipitation. DNA was purified from endotoxin by repeated extraction with triton x-114 (Sigma Chemical Co., St. Louis, MO) and tested for endotoxin using the LAL-assay (LAL-assay BioWhittaker, Walkersville, MD; lower detection limit 0.1 EU/ml) and the high sensitivity assay for endotoxin described earlier (lower detection limit 0.0014 EU/ml) (Hartmann G., and Krieg A. M. 1999. CpG DNA and LPS induce distinct patterns of activation in human monocytes. *Gene Therapy* 6:893). Endotoxin content of DNA samples was below 0.0014 U/ml. *E. coli* and calf thymus DNA were made single stranded before use by boiling for 10 minutes, followed by cooling on ice for 5 minutes. DNA samples were diluted in TE-buffer using pyrogen-free reagents.

Table H: Oligonucleotide panel used<sup>1</sup>

- 138 -

	Name (SEQ ID NO)	Sequence 5' to 3'
Starting sequence	PE 2079 (320)	<u>TCG</u> <u>ACG</u> TTC CCC CCC CCC CC
Middle base	PE 2100 (341)	TCG <b>GCG</b> TTC CCC CCC CCC CC
	PE 2082 (323)	TCG <b>CCG</b> TTC CCC CCC CCC CC
<b>Human CpG motif</b>	<b>PE 2080 (321)</b>	<b>TCG TCG</b> TTC CCC CCC CCC CC
5' flanking base	PE 2105 (346)	<b>GCG</b> TCG TTC CCC CCC CCC CC
	PE 2107 (348)	<b>ACG</b> TCG TTC CCC CCC CCC CC
	PE 2104 (345)	<b>CCG</b> TCG TTC CCC CCC CCC CC
3' flanking base	PE 2098 (339)	TCG TCG <b>CTC</b> CCC CCC CCC CC
	PE 2099 (340)	TCG TCG <b>GTC</b> CCC CCC CCC CC
	PE 2083 (324)	TCG TCG <b>ATC</b> CCC CCC CCC CC
First CpG deleted	PE 2108 (349)	CTG <u>TCG</u> TTC CCC CCC CCC CC
Second CpG deleted	PE 2106 (347)	<u>TCG</u> TCA TTC CCC CCC CCC CC
Methylation	PE 2095 (336)	<b>TZG</b> <b>TZG</b> TTC CCC CCC CCC CC
	PE 2094 (335)	TCG TCG TTC CCC CCC <b>ZCC</b> CC
Non-CpG control of 2080	PE 2078 (319)	<b>TGC</b> <b>TGC</b> TTC CCC CCC CCC CC
	PE 2101 (342)	<b>GGC</b> <b>CTT</b> TTC CCC CCC CCC CC
PS form of 2080	PS 2116 (357)	<u>TCG</u> <u>TCG</u> TTC CCC CCC CCC CC
Additional CpG motifs	PE 2059 (300)	<u>TCG</u> <u>TCG</u> TTT TGT <u>CGT</u> TTT <u>GTC</u> <u>GTT</u>
Best PS	PS 2006 (246)	<u>TCG</u> <u>TCG</u> TTT TGT <u>CGT</u> TTT <u>GTC</u> <u>GTT</u>
Methylated 2006	PS 2117 (358)	<b>TZG</b> <b>TZG</b> TTT TGT <b>ZGT</b> TTT <b>GTZ</b> <b>GTT</b>

<sup>1</sup>PE, phosphodiester; PS, phosphorothioate; bold, base exchange; bold Z, methylated cytidine; underlined, CpG dinucleotides.



- 139 -

**Cell preparation and cell culture:** Human peripheral blood mononuclear cells (PBMC) were isolated from peripheral blood of healthy volunteers by Ficoll-Paque density gradient centrifugation (Histopaque-1077, Sigma Chemical Co., St. Louis, MO) as described (Hartmann G., et al 1996 *Antisense Nucleic Acid Drug Dev* 6:291)). Cells were suspended in RPMI 1640 culture medium supplemented with 10 % (v/v) heat-inactivated (56°C, 1 h) FCS (HyClone, Logan, UT), 1.5 mM L-glutamine, 100 U/ml penicillin and 100 µg/ml streptomycin (all from Gibco BRL, Grand Island, NY) (complete medium). All compounds were purchased endotoxin-tested. Viability was determined before and after incubation with ODN by trypan blue exclusion (conventional microscopy) or by propidium iodide exclusion (flow cytometric analysis). In all experiments, 96 % to 99 % of PBMC were viable. Cells (final concentration  $1 \times 10^6$  cells/ml) were cultured in complete medium in a 5 % CO<sub>2</sub> humidified incubator at 37°C. Different oligonucleotides (see table I, concentration as indicated in the figure legends), LPS (from salmonella typhimurium, Sigma Chemical Co., St. Louis, MO) or anti-IgM were used as stimuli. Chloroquine (5 µg/ml; Sigma Chemical Co., St. Louis, MO) was used to block endosomal maturation/acidification. At the indicated time points, cells were harvested for flow cytometry as described below.

For signal transduction studies, human primary B-cells were isolated by immunomagnetic cell sorting using the VARIOMACS technique (Miltenyi Biotec Inc., Auburn, CA) as described by the manufacturer. In brief, PBMC obtained from buffy coats of healthy blood donors (Elmer L. DeGowin Blood Center, University of Iowa) were incubated with a microbeads-conjugated antibody to CD19 and passed over a positive selection column. Purity of B-cells was higher than 95%. After stimulation, whole cellular extracts (Western blot) and nuclear extracts (EMSA) for signal transduction studies were prepared.

For CpG binding protein studies, Ramos cells (human Burkitt lymphoma B cell line, ATCC CRL-1923 or CRL-1596; Intervirology 5: 319-334, 1975) were grown in complete medium. Untreated cells were harvested and cytosolic protein extracts were prepared and analyzed for the presence of CpG oligonucleotide binding proteins by EMSA and UV-crosslink as described below.

**Flow cytometry:** Staining of surface antigens was performed as previously described (Hartmann G. et al. 1998 *J Pharmacol Exp Ther* 285:920). Monoclonal

- 140 -

antibodies to HLA-DR were purchased from Immunotech, Marseille, France. All other antibodies were purchased from Pharmingen, San Diego, CA: mABs to CD19 (B43), CD40 (5C3), CD54 (HA58), CD86 (2331 (FUN-1)). IgG<sub>1,κ</sub> (MOPC-21) and IgG<sub>2b,κ</sub> were used to control for specific staining. Intracellular cytokine staining for IL-6 was performed as described (Hartmann G., and Krieg A. M. 1999. CpG DNA and LPS induce distinct patterns of activation in human monocytes. *Gene Therapy* 6:893). In brief, PBMC (final concentration  $1 \times 10^6$  cells/ml) were incubated in the presence of brefeldin A (final concentration 1  $\mu$ g/ml, Sigma Chemical Co., St. Louis, MO). After incubation, cells were harvested and stained using a FITC-labeled mAB to CD19 (B43), a PE-labeled rat anti-human IL-6 mAb (MQ2-6A3, Pharmingen) and the Fix and Perm Kit (Caltag Laboratories, Burlingame, CA). Flow cytometric data of 5000 cells per sample were acquired on a FACScan (Beckton Dickinson Immunocytometry Systems, San Jose, CA). Non-viable cells were excluded from analysis by propidium iodide staining (2  $\mu$ g/ml). Data were analyzed using the computer program FlowJo (version 2.5.1, Tree Star, Inc., Stanford, CA).

**Proliferation assay:** CFSE (5-(and-6-) carboxyfluorescein diacetate succinimidyl ester, Molecular Probes, USA) is a fluorescein-derived intracellular fluorescent label which is divided equally between daughter cells upon cell division. Staining of cells with CFSE allows both quantification and immunophenotyping (phycoerythrin-labeled antibodies) of proliferating cells in a mixed cell suspension. Briefly, PBMC were washed twice in PBS, resuspended in PBS containing CFSE at a final concentration of 5  $\mu$ M, and incubated at 37°C for 10 minutes. Cells were washed three times with PBS and incubated for five days as indicated in the figure legends. Proliferating CD19-positive B-cells were identified by decreased CFSE content using flow cytometry.

**Preparation of whole cell, nuclear and cytosolic protein extracts:** For Western blot analysis, whole cell extracts were prepared. Primary B-cells were treated with medium, the phosphodiester oligonucleotides 2080 (SEQ ID NO.: 321) or 2078 (SEQ ID NO.: 319) at 30  $\mu$ g/ml, or anti-IgM (10  $\mu$ g/ml). Cells were harvested, washed twice with ice-cold PBS containing 1 mM Na<sub>3</sub>VO<sub>4</sub>, resuspended in lysis buffer (150 mM NaCl, 10 mM TRIS pH 7.4, 1 % NP40, 1 mM Na<sub>3</sub>VO<sub>4</sub>, 50 mM NaF, 30 mg/ml leupeptin, 50 mg/ml aprotinin, 5 mg/ml antipain, 5 mg/ml pepstatin, 50  $\mu$ g/ml

- 141 -

phenylmethylsulfonylfluoride (PMSF)), incubated for 15 min on ice and spun at 14000 rpm for 10 min. The supernatant was frozen at -80 C. For the preparation of nuclear extracts, primary B-cells were resuspended in hypotonic buffer (10 mM HEPES/KOH (pH 7.9), 10 mM KCl, 0.05 % NP40, 1.5 mM MgCl<sub>2</sub>, 0.5 mM dithiothreitol (DTT), 0.5 mM PMSF, 30 mg/ml leupeptin, 50 mg/ml aprotinin, 5 mg/ml antipain, 5 mg/ml pepstatin). After 15 minutes incubation on ice, the suspension was centrifuged at 1000 x g for 5 minutes. The pelleted nuclei were resuspended in extraction buffer (20 mM HEPES (pH 7.9), 450 mM NaCl, 50 mM NaF, 20% glycerol, 1 mM EDTA, 1 mM EGTA, 1 mM DTT, 1 mM PMSF, 30 mg/ml leupeptin, 50 mg/ml aprotinin, 5 mg/ml antipain, 5 mg/ml pepstatin) and incubated on ice for one hour. The nuclear suspension was centrifuged for 10 minutes at 16,000 g at 4°C. Supernatant was collected and stored at -80°C. Cytosolic extracts for the CpG binding protein studies were prepared from unstimulated Ramos cells, which were lysed with hypotonic buffer as described for the preparation of the nuclear extract. After centrifugation, the supernatant was removed as cytoplasmic fraction and stored at -80°C. Protein concentrations were measured using a Bradford protein assay (Bio-Rad, Hercules, CA) according to the manufacturer.

**Western blot analysis:** Equal concentrations of whole cell protein extracts (25 µg/lane) were boiled in SDS sample buffer (50 mM Tris-Cl, pH 6.8; 1% β-mercaptoethanol; 2% SDS; 0.1% bromphenolblue; 10% glycerol) for 4 min before being subjected to electrophoresis on a 10 % polyacrylamide gel containing 0.1 % SDS (SDS-PAGE). After electrophoresis, proteins were transferred to Immobilon-P transfer membranes (Millipore Corp. Bedford, MA). Blots were blocked with 5 % nonfat dry milk. Specific antibodies against the phosphorylated form of extracellular receptor kinase (ERK), c-jun NH2-terminal kinase (JNK), p38 and activating transcription factor-2 (ATF-2) were used (New England BioLabs, Beverly, MA). Blots were developed in enhanced chemiluminescence reagent (ECL; Amersham International, Aylesbury, U.K.) according to the manufacturer's recommended procedure.

**Electrophoretic mobility shift assay (EMSA):** To detect the DNA-binding activity of the transcription factor activator protein-1 (AP-1) and NFκB, nuclear extracts (1 µg/lane) were analyzed by EMSA using the dsODNs 5' GAT CTA GTG ATG AGT CAG CCG GAT C 3' (SEQ ID NO.: 838) containing the AP-1 binding sequence, and the NFκB URE from the c-myc promotor region 5' TGC AGG AAG TCC GGG TTT TCC

- 142 -

CCA ACC CCC C 3' (SEQ ID NO.: 1142), as probes. ODNs were end labeled with T4-polynucleotide kinase (New England Biolabs) and ( $\gamma$ -<sup>32</sup>P) ATP (Amersham, Arlington Heights, IL). Binding reactions were performed with 1  $\mu$ g nuclear protein extract in DNA-binding buffer (10 mM Tris-HCl (pH 7.5), 40 mM MgCl<sub>2</sub>, 20 mM EDTA, 1 mM dithiothreitol, 8% glycerol and 100 - 400 ng of poly (dI-dC) with 20.000 - 40.000 cpm labeled ODN in 10  $\mu$ l total volume. Specificity of the NF $\kappa$ B bands was confirmed by competition studies with cold oligonucleotides from unrelated transcription factor binding sites (10 - 100 ng). For the supershift assay, 2  $\mu$ g of specific antibodies for c-Rel, p50 and p65 (Santa Cruz Biotechnology, Inc., Santa Cruz, CA) were added into the reaction mixture for 30 min before the radiolabeled probe was added. Following incubation for 30 minutes at room temperature loading buffer was added and the probes were electrophoresed on a 6 % polyacrylamide gel in Tris-borate-EDTA running buffer (90 mM Tris, 90 mM boric acid, 2 mM EDTA, pH 8.0). Gels were dried and then autoradiographed.

**UV-crosslinking and denaturing protein electrophoresis:** Nuclear extracts were incubated with labeled phosphodiester oligonucleotide as described for the EMSA. DNA-protein complexes were crosslinked with UV-light in a Stratalinker (Stratagene) for 10 minutes. Probes were mixed with SDS-sample buffer, boiled for 10 minutes and loaded on a 7.5% SDS-PAGE. The gel was dried on Whatman paper and autoradiographed. Plotting the distance against the molecular weight of the marker proteins yielded a standard curve which was used to calculate the approximate molecular weight of the crosslinked protein-ODN complexes. The molecular weight of the oligonucleotide was subtracted from this value to give the size.

***Example 9: Identification of an Optimal CpG motif for use alone or in combination with a T-rich ODN***

Phosphorothioate oligonucleotides containing the murine CpG motif GACGTT (SEQ ID NO.: 1143) (for example 1826 (SEQ ID NO.: 69)) and used at concentrations which are active in murine B-cells (Yi A. K., Chang M., Peckham D. W., Krieg A. M., and Ashman R. F. 1998. CpG oligodeoxyribonucleotides rescue mature spleen B cells from spontaneous apoptosis and promote cell cycle entry. *J Immunol* 160:5898), have showed little or no immunostimulatory activity on human immune cells. At higher

- 143 -

concentrations this ODN was found to demonstrate some stimulatory effect on human B cells.

In earlier studies on B-cell activation in mice, it was found that a CpG-dinucleotide flanked by two 5' purines and two 3' pyrimidines and preferably the 6mer motif 5' GACGTT 3' (SEQ ID NO:1143) was optimal for a phosphodiester  
5 oligonucleotide to be active (Krieg A. M., et al. 1995 *Nature* 374:546, Yi A. K., Chang M., et al.. 1998 *J Immunol* 160:5898).

In order to identify an optimal motif for stimulation of an immune response in humans and non-rodent vertebrates we designed a series of ODN and tested their  
10 activity. First we designed a 20 mer phosphodiester oligonucleotide with a TC dinucleotide at the 5' end preceding the optimal murine CpG motif 5' GACGTT 3' (SEQ ID NO.: 1143) and followed by a poly C tail (2079: 5' TCG ACG TTC CCC CCC CCC CC 3'(SEQ ID NO.: 320)). This oligonucleotide if added to human primary B-cells under the same conditions as found to be optimal for E. coli DNA (repeated addition at 0  
15 hours, 4 hours and 18 hours; 30 µg/ml for each time point) stimulated high levels of CD86 expression on human primary B-cells after two days. To determine the structure-function relationship of the CpG motifs, we replaced the bases adjacent to the CpG dinucleotides while maintaining the two CpG dinucleotides within the sequence. Exchange of the adenine located between both CpG dinucleotides by thymidine (2080  
20 (SEQ ID NO.: 321)) resulted in slightly higher activity. Replacement by guanosine (2100 (SEQ ID NO.: 341)) or cytidine (2082 (SEQ ID NO.: 323)) at this position showed no major changes compared to 2079 (SEQ ID NO.: 320). In contrast, replacement of the thymidine 3' to the second CpG dinucleotide by the purines guanosine (2099 (SEQ ID NO.: 340)) or adenine (2083 (SEQ ID NO.: 324)) resulted in a major drop in activity of  
25 the oligonucleotide, while the pyrimidine cytidine caused only a minor decrease. The thymidine immediately 5' to the first CpG dinucleotide was also important. Replacement of the thymidine by any other base (2105 (SEQ ID NO.: 346), guanosine; 2107 (SEQ ID NO.: 348), adenine; 2104 (SEQ ID NO.: 345), cytidine) led to a marked decrease in activity of the oligonucleotide. Elimination of the first (2108 (SEQ ID NO.:  
30 349)) or the second (2106 (SEQ ID NO.: 347)) CpG dinucleotide also partially reduced the activity.

- 144 -

The addition of more 5' GTCGTT 3' (SEQ ID NO.: 1144) CpG motifs to the phosphodiester oligonucleotide containing the 8mer duplex CpG motif (5' TCGTCGTT 3' (SEQ ID NO.: 1145), 2080 (SEQ ID NO.: 321)) did not further enhance CD86 expression on B-cells (2059 (SEQ ID NO.: 300)). An oligonucleotide with the same  
5 sequence as 2080 (SEQ ID NO.: 321) but with a phosphorothioate backbone showed no activity above background (2116 (SEQ ID NO.: 357)). This was surprising since the phosphorothioate backbone has been reported to greatly stabilize oligonucleotides and enhance CpG-induced stimulation (Krieg A. M., Yi A. K., Matson S., Waldschmidt T. J., Bishop G. A., Teasdale R., Koretzky G. A., and Klinman D. M. 1995. CpG motifs in  
10 bacterial DNA trigger direct B-cell activation. *Nature* 374:546). We therefore performed further structure-function analysis of phosphorothioate oligonucleotides containing the 5' GTCGTT 3' (SEQ ID NO.: 1144) and 5' TCGTCGTT 3' (SEQ ID NO.: 1145) motifs, which showed that additional CpG motifs (2006 (SEQ ID NO.: 246)) tended to increase the activity of phosphorothioate oligonucleotides.

15 Purified B-cells isolated from peripheral blood by immunomagnetic cell sorting were activated by CpG DNA to the same extent as unpurified B-cells within PBMC. Thus, activation of B-cells is a primary response and not a secondary effect caused by cytokines secreted by other cells.

In addition to the co-stimulatory molecule CD86, the functional stage of B-cells  
20 is characterized by other surface markers. For example, activated T helper cells stimulate B-cells by CD40 ligation, the intercellular adhesion molecule-1 (ICAM-1, CD54) mediates binding to other immune cells, and major histocompatibility complex II (MHC II) is responsible for antigen presentation. We found that B cell expression of CD40, CD54 and MHC II was upregulated by the CpG oligonucleotide 2080 (SEQ ID NO.:  
25 321). The non-CpG control oligonucleotide 2078 (SEQ ID NO.: 319) showed no activity compared to medium alone.

When PBMC were incubated for 5 days in the presence of 2080 (SEQ ID NO.: 321) (added at 0 hours, 4 hours, 18 hours and every subsequent morning), it was intriguing that a subpopulation of lymphocytes increased in cell size (FSC) and became  
30 more granular (SSC). To examine if this subpopulation represented proliferating B-cells, we stained freshly isolated PBMC with CFSE (5-(and-6-) carboxyfluorescein diacetate succinimidyl ester) at day 0 and incubated them for 5 days with 2080 (SEQ ID NO.: 321)

- 145 -

as above. CFSE is a fluorescent molecule that binds irreversibly to cell proteins. Each cell division decreases CFSE stain by 50 %. Cells staining low with CFSE (proliferating cells) were found to be mainly CD19-positive B-cells. The oligonucleotide 2080 (SEQ ID NO.: 321) induced 60 to 70 % of CD19 positive B-cells to proliferate within 5 days.

5 The control oligonucleotide 2078 (SEQ ID NO.: 319) induced less than 5 % of B-cells to proliferate. Proliferating B-cells (CFSE low) showed a larger cell size (FSC) and higher granularity.

Proliferating B-cells expressed higher levels of CD86 than non-proliferating cells (not shown). In agreement with this finding, the oligonucleotide panel tested above for  
10 induction of CD86 expression resulted in an almost identical pattern of B-cell proliferation. Replacement of the 3' thymidine reduced activity more than changing the thymidine in the middle position.

***Example 10: B-cell activation requires endosomal maturation/acidification***

15 It has previously been shown that chloroquine, an inhibitor of endosomal acidification, blocks CpG-mediated stimulation of murine antigen presenting cells and B-cells, while not influencing LPS-mediated effects (Hacker H., et al 1998 *Embo J* 17:6230, Yi A. K. et al 1998 *J Immunol* 160:4755, Macfarlane D. E., and Manzel L. 1998 *J Immunol* 160:1122). We found that the addition of 5 µg/ml chloroquine  
20 completely blocked CpG DNA-mediated induction of CD86 expression on primary B-cells (MFI CD86: 2006 (SEQ ID NO.: 246), 4.7 vs 1.4; E. coli DNA, 3.4 vs. 1.4; medium only, 0.9; n=4). Furthermore, chloroquine completely inhibited the induction of B-cell proliferation by the phosphorothioate oligonucleotide 2006 (SEQ ID NO.: 246) measured with the CFSE proliferation assay as well as with the standard . These results  
25 suggest that as with murine cells, activation of human B-cells by CpG DNA requires the uptake of DNA in endosomes and subsequent endosomal acidification.

***Example 11: Analysis of sub-cellular events resulting upon human B cell stimulation with optimal human ODN.***

30 Since the CpG motif requirement for maximal B-cell activation is substantially different between mouse (GACGTT) (SEQ ID NO:1143) and humans (TCGTCGTT) (SEQ ID NO:1145), we were interested if the basic intracellular signaling events are

- 146 -

comparable. Rapid induction of NF $\kappa$ B binding activity has been found earlier in murine B-cells and macrophages (Stacey K. J., et al 1996 *J Immunol* 157:2116, Yi A. K et al 1998 *J Immunol* 160:4755). To investigate the NF $\kappa$ B response to CpG DNA in humans, human primary B-cells were isolated from peripheral blood by immunomagnetic cell sorting and incubated with the CpG oligonucleotide 2080 (SEQ ID NO.: 321), the non-CpG control oligonucleotide 2078 (SEQ ID NO.: 319), or medium. At the indicated time points, cells were harvested and nuclear extracts were prepared. In the presence of CpG oligonucleotide, NF $\kappa$ B binding activity was increased within one hour and maintained up to 18 hours (latest time point examined). The non-CpG control oligonucleotide 2078 (SEQ ID NO.: 319) did not show enhanced NF $\kappa$ B activity compared to cells incubated with medium only. The NF $\kappa$ B band was identified by cold competition, and shown to consist of p50 and p65 subunits by supershift assay.

The activating protein-1 (AP-1) transcription factor is involved in the regulation of immediate early genes and cytokine expression (Karin M. 1995. The regulation of AP-1 activity by mitogen-activated protein kinases. *J Biol Chem* 270:16483). In murine B-cells, AP-1 binding activity is induced in response to CpG DNA (Yi A. K., and Krieg A. M. 1998. Rapid induction of mitogen-activated protein kinases by immune stimulatory CpG DNA. *J Immunol* 161:4493). To determine whether this transcription factor would also be induced by CpG DNA in humans, we examined AP-1 DNA binding activity in human primary B-cells. Cells were incubated with the CpG oligonucleotide 2080 (SEQ ID NO.: 321) or the control oligonucleotide 2078 (SEQ ID NO.: 319). Nuclear extracts were prepared and the AP-1 binding activity was analyzed by EMSA. AP-1 binding activity was enhanced within one hour, and increased up to 18 hours (latest time point examined), showing a sustained response.

Since AP-1 activity is induced by many stimuli (Angel P., and Karin M. 1991. The role of Jun, Fos and the AP-1 complex in cell-proliferation and transformation. *Biochim Biophys Acta* 1072:129), we were interested in signal transduction pathways upstream of AP-1. The AP-1 transcription factor complex integrates different mitogen activated protein kinase (MAPK) pathways (Karin M. 1995. The regulation of AP-1 activity by mitogen-activated protein kinases. *J Biol Chem* 270:16483). Western blots were performed using whole cell extracts from primary B-cells incubated with the CpG oligonucleotide 2080 (SEQ ID NO.: 321), the control 2078 (SEQ ID NO.: 319), or



- 147 -

medium only. Specific antibodies to the phosphorylated form of JNK, p38, ATF-2 and ERK were used. Strong induction of JNK phosphorylation was found 30 min and 60 min after exposure to CpG-DNA, while the non-CpG oligonucleotide showed no activity above background. The protein kinase p38, another stress activated protein kinase (SAPK), was also phosphorylated in response to CpG DNA within 60 min. ATF-2, a substrate of both p38 and JNK (Gupta S., Campbell D., Derijard B., and Davis R. J. 1995. Transcription factor ATF2 regulation by the JNK signal transduction pathway. *Science* 267:389) and a component of the AP-1 complex, showed weak phosphorylation after 30 min which increased after 60 min. CpG DNA failed to induce substantial phosphorylation of ERK. In contrast, anti-IgM, stimulating the B-cell receptor, did trigger phosphorylation of ERK. Anti-IgM activated different isoforms of JNK than CpG DNA.

***Example 12: Assay for in vivo adjuvant activity.***

An *in vitro* screening assay to identify ODN useful as an adjuvant *in vivo* in humans and other non-rodent animals was developed. Since we saw not only quantitative but also qualitative differences in activities of different CpG ODN in mice, we first screened a panel of CpG and non-CpG control ODN on mouse cells to find *in vitro* assays with reliable and strong correlation to *in vivo* adjuvant activity with hepatitis B surface antigen (HBsAg). We then systematically tested a panel of more than 250 ODN in corresponding human assays to identify sequences with *in vitro* immunostimulatory activity. We next examined if the ODN with the highest activity in these human assays also activate B cell proliferation in chimpanzees and monkeys, and finally, if they are active as adjuvants with HBsAg in chimpanzees and cynomolgus monkeys *in vivo*. These studies revealed that the sequence, number and spacing of individual CpG motifs contribute to the immunostimulatory activity of a CpG phosphorothioate ODN. An ODN with a TC dinucleotide at the 5' end followed by three 6mer CpG motifs (5' GTCGTT 3') separated by TT dinucleotides consistently showed the highest activity for human, chimpanzee, and rhesus monkey leukocytes. Chimpanzees or monkeys vaccinated once against hepatitis B with this CpG ODN adjuvant developed 15 times higher anti-HBs antibody titers than those receiving vaccine alone.

### ***Materials and Methods***

**Oligodeoxynucleotides:** Phosphorothioate-modified ODN were purchased from Operon Technologies (Alameda, CA) and Hybridon Specialty Products (Milford, MA). ODN were tested for endotoxin using the LAL-assay (LAL-assay BioWhittaker, Walkersville, MD; lower detection limit 0.1 EU/ml). For *in vitro* assays, ODN were diluted in TE-buffer (10 mM Tris, pH 7.0, 1 mM EDTA), and stored at -20° C. For *in vivo* use, ODN were diluted in phosphate buffered saline (0.1 M PBS, pH 7.3) and stored at 4°C. All dilutions were carried out using pyrogen-free reagents.

**Mouse spleen cell cultures:** Spleens were removed from 6-12 week old female BALB/c (The Jackson Laboratory),  $2 \times 10^6$  splenocytes were cultured with 0.2  $\mu$ M ODN for 4 hours (TNF- $\alpha$ ) or 24 hours (IL-6, IFN- $\gamma$ , IL-12), and cytokines were detected by ELISA as previously described (Yi A. K., Klinman D. M., Martin T. L., Matson S., and Krieg A. M. 1996. Rapid immune activation by CpG motifs in bacterial DNA. Systemic induction of IL-6 transcription through an antioxidant-sensitive pathway. *J Immunol* 157:5394). To evaluate CpG-induced B cell proliferation, spleen cells were depleted of T cells with anti-Thy-1.2 and complement and centrifugation over lympholyte M<sup>®</sup> (Cedarlane Laboratories, Hornby, ON, Canada), cultured for 44 hours with the indicated ODN, and then pulsed for 4 hours with 1  $\mu$ Ci of <sup>3</sup>H thymidine as described previously (Krieg A. M., Yi A. K., Matson S., Waldschmidt T. J., Bishop G. A., Teasdale R., Koretzky G. A., and Klinman D. M. 1995. CpG motifs in bacterial DNA trigger direct B-cell activation. *Nature* 374:546). To examine NK cell lytic activity murine spleen cells were depleted of B cells using magnetic beads coated with goat anti-mouse Ig as previously detailed (Ballas Z. K., and Rasmussen W. 1993. Lymphokine-activated killer cells. VII. IL-4 induces an NK1.1<sup>+</sup>CD8  $\alpha^+\beta^-$  TCR- $\alpha\beta$  B220<sup>+</sup> lymphokine-activated killer subset. *J Immunol* 150:17). Cells were cultured at  $5 \times 10^6$ /well in 24-well plates and harvested at 18 hours for use as effector cells in a standard 4 hour <sup>51</sup>Cr-release assay against YAC-1 target cells. One unit (LU) was defined as the number of cells needed to effect 30 % specific lysis.

**Immunization of mice against HBsAg and evaluation of the humoral response:** Groups of 6-8 week old female BALB/c mice (n = 5 or 10, Charles River, Montreal, QC) were immunized against HBsAg as previously described (Davis H. L., et

- 149 -

al 1998 *J Immunol* 160:870). In brief, each mouse received a single IM injection of 50  $\mu$ l PBS containing 1  $\mu$ g recombinant HBsAg (Medix Biotech, Foster City, CA) and 10  $\mu$ g of CpG ODN or non-CpG ODN as a sole adjuvant or combined with alum (Alhydrogel "85", Superfos Biosector, Vedbaek, Denmark; 25 mg  $\text{Al}^{3+}$ /mg HBsAg).

- 5 Control mice were immunized with HBsAg without adjuvant or with alum. Plasma was recovered from mice at various times after immunization and Abs specific to HBsAg (anti-HBs) were quantified by end-point dilution ELISA assay (in triplicate) as described previously (Davis H. L et al 1998 *J Immunol* 160:870). End-point titers were defined as the highest plasma dilution that resulted in an absorbance value (OD450) two times  
10 higher than that of non-immune plasma with a cut-off value of 0.05.

- Isolation of primate PBMC and cell culture:** Peripheral blood mononuclear cells (PBMC) were isolated from peripheral blood of healthy volunteers, chimpanzees or rhesus or cynomolgus monkeys by Ficoll-hypaque density gradient centrifugation (Histopaque-1077, Sigma Chemical Co., St. Louis, MO) as described (Hartmann G., et al  
15 1996 *Antisense Nucleic Acid Drug Dev* 6:291). Cells were suspended in RPMI 1640 culture medium supplemented with 10 % (v/v) heat-inactivated ( $56^{\circ}\text{C}$ , 1 h) FCS (HyClone, Logan, UT), 1.5 mM L-glutamine, 100 U/ml penicillin and 100  $\mu\text{g}/\text{ml}$  streptomycin (all from Gibco BRL, Grand Island, NY) (complete medium). Cells (final concentration  $1 \times 10^6$  cells/ml) were cultured in complete medium in a 5 %  $\text{CO}_2$   
20 humidified incubator at  $37^{\circ}\text{C}$ . ODN and LPS (from *Salmonella typhimurium*, Sigma Chemical Co., St. Louis, MO) or anti-IgM were used as stimuli. For measurement of human NK lytic activity, PBMC were incubated at  $5 \times 10^6$ /well in 24-well plates. Cultures were harvested after 24 hours, and cells were used as effectors in a standard 4 hours  $^{51}\text{Cr}$ -release assay against K562 target cells as previously described (Ballas Z. K.,  
25 Rasmussen W. L., and Krieg A. M. 1996. Induction of NK activity in murine and human cells by CpG motifs in oligodeoxynucleotides and bacterial DNA. *J Immunol* 157:1840; Ballas Z. K., and Rasmussen W. 1993. Lymphokine-activated killer cells. VII. IL-4 induces an  $\text{NK1.1}^+\text{CD8}^+\text{TCR-}\alpha\beta^- \text{B220}^+$  lymphokine-activated killer subset. *J Immunol* 150:17). For B cell proliferation, 1  $\mu\text{Ci}$  of  $^3\text{H}$  thymidine was added 18 hours  
30 before harvest, and the amount of  $^3\text{H}$  thymidine incorporation was determined by scintillation counting at day 5. Standard deviations of the triplicate wells were < 5%.

**Flow cytometry on primate PBMC:** Surface antigens on primate PBMC were stained as previously described (Hartmann G et al 1998 *J Pharmacol Exp Ther* 285:920). Monoclonal antibodies to CD3 (UCHT1), CD14 (M5E2), CD19 (B43), CD56 (B159), CD69 (FN50) and CD86 (2331 (FUN-1)) were purchased from Pharmingen, San Diego, CA. IgG<sub>1</sub>, $\kappa$  (MOPC-21) and IgG<sub>2b</sub>, $\kappa$  (Hartmann G et al 1999 *PNAS* 96:9305) were used to control for non-specific staining. NK cells were identified by CD56 expression on CD3, CD14 and CD19 negative cells, whereas B cells were identified by expression of CD19. Flow cytometric data from 10000 cells per sample were acquired on a FACScan (Beckton Dickinson Immunocytometry Systems, San Jose, CA). The viability of cells within the FSC/SSC gate used for analysis was examined by propidium iodide staining (2  $\mu$ g/ml) and found to be higher than 98 %. Data were analyzed using the computer program FlowJo (version 2.5.1, Tree Star, Inc., Stanford, CA).

**Immunization of chimpanzees and cynomolgus monkeys against HBsAg and evaluation of the humoral response:** Fourteen cynomolgus monkeys (2.0–3.5 kg) were immunized with a pediatric dose of Engerix-B (SmithKline Beecham Biologicals, Rixensart, BE) containing 10  $\mu$ g HBsAg adsorbed to alum (25 mg Al<sup>3+</sup>/mg HBsAg). This was administered alone (n=5), or combined with CpG ODN 1968 (n=5, 500  $\mu$ g) or CpG ODN 2006 (SEQ ID NO.: 246) (n=4, 150  $\mu$ g). Four chimpanzees (10–20 kg) were immunized in the same fashion with two receiving control vaccine (Engerix-B only) and two receiving experimental vaccine (Engerix-B plus 1 mg CpG ODN 2006). All vaccines were administered IM in the right anterior thigh in a total volume of 1 ml. Monkeys were maintained in the animal facility of the Primate Research Center (Bogor, Indonesia) and chimpanzees were housed at Bioqual (Rockville, MD). Animals were monitored daily by animal care specialists. No symptoms of general ill health or local adverse reactions at the injection site were noted. Plasma was recovered by IV puncture prior to and at various times after immunization and was stored frozen (-20°C) until assayed for antibodies. Anti-HBs antibodies were detected using a commercial ELISA kit (Monolisa Anti-HBs; Sanofi-Pasteur, Montreal, QC) and titers were expressed in mIU/ml based on comparison with WHO defined standards (Monolisa Anti-HBs Standards; Sanofi-Pasteur).

## Results

- 151 -

Identification of CpG ODN with different profiles of *in vitro* immune activities:

Our studies showed that the precise bases on the 5' and 3' sides of a CpG dinucleotide within a CpG motif may have an impact on the level of immune activation of a synthetic ODN, but it has been unclear whether different CpG motifs might display different immune effects. To evaluate this possibility, we tested a panel of CpG ODN for their ability to induce NK lytic activity, B cell proliferation, and to stimulate synthesis of TNF- $\alpha$ , IL-6, IFN- $\gamma$  and IL-12 in murine spleen cells. Immunostimulatory activity of ODN without CpG motifs (ODN 1982 (SEQ ID NO.: 225), ODN 1983 (SEQ ID NO.: 226)) was negative or weak compared to CpG ODN. ODN with non optimal CpG motifs (ODN 1628 (SEQ ID NO.: 767), ODN 1758 (SEQ ID NO.: 1)) were less active than ODN containing CpG motifs flanked by two 5' purines and two 3' pyrimidines (ODN 1760 (SEQ ID NO.: 3), ODN 1826 (SEQ ID NO.: 69), ODN 1841 (SEQ ID NO.: 84)). ODN 1826 containing two optimal murine CpG motifs (5' GACGTT 3') (SEQ ID NO.: 1143) had the highest activity for 5 of 6 measured endpoints. Except for ODN 1628, all ODN showed a generally similar pattern of activity (NK cell-mediated lysis, B cell proliferation, IL-12, IL-6, TNF  $\alpha$ , IFN- $\gamma$ ). Of note, ODN 1628, which was unique in this panel for containing two G-rich regions, showed preferential induction of IFN- $\gamma$  synthesis but relatively low stimulation of the other activities.

Identification of *in vitro* assays which correlate with *in vivo* adjuvant activity:

Since adjuvant activity is an *in vivo* endpoint, we were interested in identifying *in vitro* assays that would predict the adjuvant activity of a CpG ODN *in vivo*. The same ODN used for *in vitro* endpoints therefore were tested for their adjuvant activity to immunize mice against HBsAg. This was carried out both with ODN alone and with ODN combined with alum, since earlier studies had shown strong synergy for CpG ODN and alum adjuvants (PCT Published Patent Application WO98/40100).

BALB/c mice immunized with HBsAg without adjuvant attained only low titers of anti-HBs by 4 weeks, and this was not affected by addition of control ODN. In contrast, addition of CpG ODN raised anti-HBs titers by 5 to 40 fold, depending on the sequence used. When alum was added, titers of anti HBs were approximately 6 times higher than with HBsAg alone. Specifically, control ODN had no effect and the various CpG ODN augmented these titers 2 to 36 fold. Results obtained with the different ODN

- 152 -

alone correlated very strongly ( $r = 0.96$ ) with those obtained using the same ODN plus alum. When linear regression was performed, a very high degree of correlation was found between certain *in vitro* assays and *in vivo* augmentation of anti-HBs titers. Of all the *in vitro* endpoints examined, the induction of NK lytic activity showed the best  
5 correlation to *in vivo* adjuvant activity (without alum,  $r = 0.98$ ; with alum,  $r = 0.95$ ;  $p < 0.0001$ ). A good correlation regarding adjuvant activity was also obtained for B-cell stimulation ( $r = 0.84$  and  $0.7$ ), as well as secretion of TNF- $\alpha$  ( $r = 0.9$  and  $0.88$ ), IL-12 ( $r = 0.88$  and  $0.86$ ) and IL-6 ( $r = 0.85$  and  $0.91$ ). The one *in vitro* assay that did not correlate well with the *in vivo* results was IFN- $\gamma$  secretion ( $r = 0.57$  and  $0.68$ ). These data  
10 demonstrate that *in vitro* assays for NK lytic activity, B cell activation and production of TNF- $\alpha$ , IL-6 and IL-12 provide valuable information *in vitro* to predict the adjuvant activity of a given ODN *in vivo*.

Screening of a phosphorothioate ODN panel to activate human NK cells: In previous studies we found that synthesis of inflammatory cytokines by human PBMC is  
15 induced by extremely low amounts of endotoxin (induced TNF- $\alpha$  secretion is detectable with just 6 pg/ml endotoxin, 2 logs more sensitive than murine immune cells). In contrast, activation of human B cells and induction of human NK cell lytic activity with endotoxin is low even at high endotoxin concentrations. Based on these results we selected activation of NK cells (lytic activity and CD69 expression) and B cells  
20 (proliferation and CD86 expression) as the most highly specific and reproducible assays with low inter-subject variability and used these assays for *in vitro* screening of a pool of ODN.

First we studied the effect of phosphorothioate ODN containing various combinations and permutations of CpG motifs on NK cell-mediated lysis of target cells.  
25 For clarity and ease of presentation, only data with selected representative CpG and control ODN are shown. Human PBMC were incubated with different phosphorothioate ODN (6  $\mu\text{g/ml}$ ) for 24 hours and tested for their ability to lyse  $^{51}\text{Cr}$ -labeled K562 cells. ODN with two 6-mer CpG motifs (either 5' GACGTT 3' (SEQ ID NO.: 1143) or 5' GTCGTT 3' (SEQ ID NO.: 1144)) in combination with a TpC at the 5' end of the ODN  
30 (ODN 1840 5' TCCATGTCGTTCTGTCGTT 3' (SEQ ID NO.: 83), ODN 1851 5' TCCTGACGTTCTGACGTT 3' (SEQ ID NO.: 94) or with at least three 6-mer motifs without a TpC at the 5' end (ODN 2013 (SEQ ID NO.: 253)) show intermediate activity.

- 153 -

High activity was found when the 5' TpC directly preceded a 6-mer human CpG motif (5' TCGTCGTT 3' (SEQ ID NO.: 1145)) and was followed by two 6-mer motifs (ODN 2005 (SEQ ID NO.: 245), ODN 2006 (SEQ ID NO.: 246) and ODN 2007 (SEQ ID NO.: 247)). The best results were obtained when the 6-mer CpG motifs were separated from  
5 each other and from the 5' 8-mer CpG motif by TpT (ODN 2006 (SEQ ID NO.: 246)).

Expression of the activation marker CD69 is rapidly upregulated on the surface of NK cells subsequent to stimulation. To confirm the results from the NK cell lysis assay, PBMC were incubated for 18 hours with ODN (2 µg/ml). CD69 expression was examined on CD56 positive NK cells (CD3, CD14 and CD19 negative). Although  
10 induction of CD69 expression was less sequence restricted than stimulation of NK cell functional activity, control ODN (ODN 1982, ODN 2116, ODN 2117, ODN 2010) showed only low activity similar to background levels. ODN with two human CpG motifs separated by 5' TTTT 3' (ODN 1965 (SEQ ID NO.: 208)) or four human CpG motifs without spacing (ODN 2013 (SEQ ID NO.: 253)) were relatively more active at  
15 inducing CD69 expression than at stimulating NK cell lytic activity. Optimal NK cell functional activity, as well as CD69 expression, was obtained with ODNs containing a TpC dinucleotide preceding the human CpG motif, and additional human motifs within the sequence (ODN 2006 (SEQ ID NO.: 246), ODN 2007 (SEQ ID NO.: 247)).

Activity of phosphorothioate ODN for stimulating human B cells: In preliminary  
20 experiments we found that the percentage of proliferating B cells (CFSE assay, see methods section) correlated with the surface expression of the co-stimulatory CD86 on B cells, as measured by flow cytometry. Thus we used CD86 expression on B cells to screen a panel of ODN for their immunostimulatory activity. PBMC were incubated with 0.6 µg/ml ODN. Expression of CD86 (mean fluorescence intensity, MFI) was examined  
25 on CD19 positive B cells. A poly C ODN (ODN 2017 (SEQ ID NO.: 257)) or ODN without CpG dinucleotides (ODN 1982 (SEQ ID NO.: 225)) failed to stimulate human B cells under these experimental conditions. A phosphorothioate ODN (ODN 2116 (SEQ ID NO.: 256)) with one optimal human CpG motif preceded by a TpC (5' TCGTCGTT 3' (SEQ ID NO.: 1145)) had low activity. The presence of one human 6-mer CpG motif  
30 (5' GTCGTT 3' (SEQ ID NO.: 1144)) had no activating effect. Two of these CpG motifs within the sequence showed no (ODN 1960 (SEQ ID NO.: 203), ODN 2016 (SEQ ID NO.: 256)) or intermediate (ODN 1965 (SEQ ID NO.: 208)) activity dependent on the

- 154 -

sequence context. If the ODN was composed of three or four copies of this motif (ODN 2012 (SEQ ID NO.: 252), ODN 2013 (SEQ ID NO.: 253), ODN 2014 (SEQ ID NO.: 254)), intermediate activity on B cells could be detected. The combination of the human 8-mer CpG motif on the 5' end of the ODN with two 6-mer CpG motifs (ODN 2005 (SEQ ID NO.: 245), ODN 2006 (SEQ ID NO.: 246), ODN 2007 (SEQ ID NO.: 247), ODN 2102 (SEQ ID NO.: 343), ODN 2103 (SEQ ID NO.: 344)) led to a considerable increase in the ability of the ODN to stimulate B cells. The spacing between the single motifs was critical. The separation of CpG motifs by TpT was preferable (ODN 2006 (SEQ ID NO.: 246)) compared to unseparated CpG motifs (ODN 2005 (SEQ ID NO.: )); also compare ODN 1965 (SEQ ID NO.: 208) to ODN 1960 (SEQ ID NO.: 203)). The human 6-mer CpG motif (5' GTCGTT 3') was better than the optimal mouse 6-mer CpG motif (5' GACGTT 3' (SEQ ID NO.: 246)) when combined with the human 8-mer CpG motif on the 5' end (ODN 2006 vs. ODN 2102 (SEQ ID NO.: 343) and ODN 2103 (SEQ ID NO.: 344)). A (TCG)<sub>poly</sub> ODN was inactive or only weakly active, as were ODN containing CpG dinucleotides flanked by guanines or other CpG dinucleotides (ODN 2010 (SEQ ID NO.: 250)). Taken together, the findings for NK cells and B cells showed consistently that of the ODN tested, ODN 2006 (SEQ ID NO.: 246) has the highest immunostimulatory activity on human immune cells.

Comparative analysis of potency of CpG phosphorothioate ODNs in different primates: Different CpG motifs are optimal to activate murine and human immune cells. Furthermore, the number and location of CpG motifs within an active phosphorothioate ODN is different in mice and humans. We were interested to know if CpG phosphorothioate ODN show a similar activity among different species of primates. We compared a panel of CpG ODN for their ability to induce B cell proliferation in humans, chimpanzees and rhesus or cynomolgus monkeys. The capability of ODN to stimulate human B cell proliferation (Table J) correlated well with their ability to induce CD86 expression on B cells. ODN 2006 (SEQ ID NO.: 246), which showed the highest activity in human B cells and NK cells, was also the most active in stimulating chimpanzee and rhesus monkey B cell proliferation (Table J). ODN 1968 (SEQ ID NO.: 211) and ODN 2006 (SEQ ID NO.: 246) gave the highest activation of cynomolgus monkey B-cells *in vitro* (SI of 25 and 29 respectively at 6 µg ODN/ml). Surprisingly, CpG ODN 2007 (SEQ ID NO.: 247), which displayed similarly high activity as the optimal ODN 2006



- 155 -

(SEQ ID NO.: 246) in human cells, did not stimulate Rhesus monkey or chimpanzee B cell proliferation, and the ODN 1968 (SEQ ID NO.: 211) showed low activity. CpG ODN originally identified with high activity in mice (ODN 1760 (SEQ ID NO.: 3), ODN 1826 (SEQ ID NO.: 69)) showed little activity in monkeys (Table J).

5 **Table J:** Proliferative response of PBMC to phosphorothioate CpG ODN in primates

	Humans	Chimpanzee	Rhesus monkey
No addition	0.5+-0.1	0.5+-0.1	0.5+-0.0
ODN 1760 (SEQ ID NO.: 3)	23+-7	0.3+-0.1	0.5+-0.3
ODN 1826 (SEQ ID NO.: 69)	0.8+-0.1	0.4+-0.1	0.6+-0.1
ODN 1968 (SEQ ID NO.: 211)	35+-9	20.0+-3.8	1.9+-0.7
ODN 1982 (SEQ ID NO.: 225)	9.7+-1.1	2.5+-1.1	0.7+-0.1
<b>ODN 2006</b> (SEQ ID NO.: 246)	<b>58+-8</b>	<b>27.4+-8.9</b>	<b>6.3+-3.3</b>
ODN 2007 (SEQ ID NO.: 247)	47+-11	0.5+-0.1	0.4+-0.2

PBMC were prepared from peripheral blood and incubated with ODN (0.6 µg/ml) as indicated for five days. Proliferation was measured by uptake of <sup>3</sup>H/thymidine (cpm/1000) during the last 18 hours. More than 95 % of proliferating cells were B-cells as determined using the CFSE assay. Four human probands, six chimpanzees and two rhesus monkeys were tested.

In vivo adjuvant activity of CpG ODN in chimpanzees and cynomolgus monkeys: In order to evaluate whether CpG ODN with strong *in vitro* stimulatory effects on primate cells had detectable adjuvant activity *in vivo*, Cynomolgus monkeys and chimpanzees were immunized with Engerix B, which comprises HBsAg adsorbed to

- 156 -

alum, alone or with added ODN 1968 (500 µg) or ODN 2006 (SEQ ID NO.: 246) (1 mg) respectively. Compared to controls not receiving CpG ODN, anti-HBs titers at 4 weeks post-prime and 2 weeks post-boost were 66- and 16-fold higher respectively in the monkeys, and 15- and 3-fold higher in the chimpanzees (Table K). Thus a clear adjuvant effect of CpG ODN was seen, and this was particularly striking after a single immunization.

**Table K** Anti-HBs responses in primates immunized against HBsAg with CpG ODN<sup>3</sup>

<u>Primate species</u>	<u>n</u>	<u>CpG ODN</u>	<u>Anti-HBs (mIU/ml)</u>	
			<u>4 wks post-prime</u>	<u>2 wks post-boost</u>
Cynomolgus monkey	5	None	15 ± 44	4880 ± 13113
	5	ODN 1968 (500 µg) (SEQ ID NO. 211)	995 ± 1309	76449 ± 42094
Chimpanzee	2	None	6, 11	3712, 4706
	2	ODN 2006 (1 mg) (SEQ ID NO. 246)	125, 135	9640, 16800

<sup>3</sup>Animals were immunized by IM injection of Engerix B containing 10 µg HBsAg adsorbed to alum, alone or with added CpG ODN. Cynomolgus monkeys were boosted at 10 wks and chimpanzees were boosted at 4 wks post-prime. Anti-HBs was determined by ELISA assay; values for monkeys are GMT ± SEM (n=5) whereas individual values for the two chimpanzees in each group are provided.

I claim:

- 157 -

**CLAIMS**

1. A method of stimulating an immune response, comprising  
administering an immunostimulatory nucleic acid selected from the group  
consisting of a Py-rich nucleic acid and a TG nucleic acid, to a non-rodent subject in an  
5 amount effective to induce an immune response in the non-rodent subject.
2. The method of claim 1, wherein the immunostimulatory nucleic acid is a T-  
rich nucleic acid.
3. The method of claim 2, wherein the T-rich immunostimulatory nucleic acid is  
a poly T nucleic acid comprising  
10 5' TTTT 3'.
4. The method of claim 3, wherein the poly T nucleic acid comprises  
5' X<sub>1</sub> X<sub>2</sub> TTTT X<sub>3</sub> X<sub>4</sub> 3'  
wherein X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub> and X<sub>4</sub> are nucleotides.
5. The method of claim 3, wherein the T-rich immunostimulatory nucleic acid  
15 comprises a plurality of poly T nucleic acid motifs.
6. The method of claim 4, wherein X<sub>1</sub>X<sub>2</sub> is TT.
7. The method of claim 4, wherein X<sub>3</sub>X<sub>4</sub> is TT.
8. The method of claim 4, wherein X<sub>1</sub>X<sub>2</sub> is selected from the group consisting of  
TA, TG, TC, AT, AA, AG, AC, CT, CC, CA, GT, GG, GA, and GC.
- 20 9. The method of claim 4, wherein X<sub>3</sub>X<sub>4</sub> is selected from the group consisting of  
TA, TG, TC, AT, AA, AG, AC, CT, CC, CA, GT, GG, GA, and GC.
10. The method of claim 3, wherein the T-rich immunostimulatory nucleic acid  
comprises a nucleotide composition of greater than 25% T.
11. The method of claim 1, wherein the T-rich immunostimulatory nucleic acid  
25 comprises a nucleotide composition of greater than 35% T.

- 158 -

12. The method of claim 1, wherein the T-rich immunostimulatory nucleic acid comprises a nucleotide composition of greater than 40% T.

13. The method of claim 1, wherein the T-rich immunostimulatory nucleic acid comprises a nucleotide composition of greater than 50% T.

5        14. The method of claim 1, wherein the T-rich immunostimulatory nucleic acid comprises a nucleotide composition of greater than 60% T.

15. The method of claim 1, wherein the T-rich immunostimulatory nucleic acid comprises a nucleotide composition of greater than 80% T.

10        16. The method of claim 1, wherein the immunostimulatory nucleic acid comprises at least 20 nucleotides.

17. The method of claim 1, wherein the immunostimulatory nucleic acid comprises at least 24 nucleotides.

18. The method of claim 1, wherein the immunostimulatory nucleic acid has a nucleotide backbone which includes at least one backbone modification.

15        19. The method of claim 18, wherein the backbone modification is a phosphorothioate modification.

20. The method of claim 18, wherein the nucleotide backbone is chimeric.

21. The method of claim 18, wherein the nucleotide backbone is entirely modified.

20        22. The method of claim 1, wherein the immunostimulatory nucleic acid is free of CpG dinucleotides.

23. The method of claim 1, wherein the immunostimulatory nucleic acid is free of unmethylated CpG dinucleotides.

25        24. The method of claim 1, wherein the immunostimulatory nucleic acid is free of methylated CpG dinucleotides.

- 159 -

25. The method of claim 1, wherein the immunostimulatory nucleic acid is free of poly-C sequences.

26. The method of claim 1, wherein the immunostimulatory nucleic acid includes a poly-A sequence.

5        27. The method of claim 20, wherein the immunostimulatory nucleic acid includes a poly-G sequence.

28. The method of claim 1, wherein the immunostimulatory nucleic acid comprises a nucleotide composition of greater than 25% C.

10       29. The method of claim 1, wherein the immunostimulatory nucleic acid comprises a nucleotide composition of greater than 25% A.

30. The method of claim 1, wherein the immunostimulatory nucleic acid is administered orally.

31. The method of claim 1, wherein the immunostimulatory nucleic acid is administered locally.

15       32. The method of claim 1, wherein the immunostimulatory nucleic acid is administered in a sustained release device.

33. The method of claim 1, wherein the immunostimulatory nucleic acid is administered mucosally to a mucosal surface.

20       34. The method of claim 33, wherein the immune response is a mucosal immune response.

35. The method of claim 33, wherein the immune response is a systemic immune response.

36. The method of claim 33, wherein the mucosal surface is selected from the group consisting of an oral, nasal, rectal, vaginal, and ocular surface.

25       37. The method of claim 1, further comprising exposing the subject to an antigen and wherein the immune response is an antigen-specific immune response.

- 160 -

38. The method of claim 37, wherein a nucleic acid vector which encodes the antigen is administered to the subject, and wherein the nucleic acid vector is separate from the immunostimulatory nucleic acid.

39. The method of claim 37, wherein the antigen is a peptide antigen.

5        40. The method of claim 1, further comprising isolating an immune cell from the subject, contacting the immune cell with an effective amount to activate the immune cell of the immunostimulatory nucleic acid and re-administering the activated immune cell to the subject.

41. The method of claim 40, wherein the immune cell is a leukocyte.

10       42. The method of claim 41, further comprising contacting the immune cell with an antigen.

43. The method of claim 40, wherein the antigen is selected from the group consisting of a tumor antigen, a viral antigen, a bacterial antigen, and a parasitic antigen.

44. The method of claim 40, wherein the immune cell is a dendritic cell.

15       45. The method of claim 1, wherein the subject has or is at risk of developing asthma and the method is a method of treating or preventing asthma in the subject.

46. The method of claim 1, wherein the subject has or is at risk of developing allergy and the method is a method of treating or preventing allergy.

20       47. The method of claim 1, wherein the subject has cancer and the method is a method of treating the cancer.

48. The method of claim 47, wherein the cancer is selected from the group consisting of biliary tract cancer; brain cancer; breast cancer; cervical cancer; choriocarcinoma; CNS cancer; colon cancer; connective tissue cancer; endometrial cancer; eye cancer; gastric cancer; intraepithelial neoplasms; larynx cancer; lymphomas; 25    Hodgkin's lymphoma; liver cancer; lung cancer (e.g. small cell and non-small cell); melanoma; neuroblastomas; oral cancer; oral cavity cancer; ovarian cancer; pancreas

- 161 -

cancer; prostate cancer; rectal cancer; sarcomas; thyroid cancer; and renal cancer, as well as other carcinomas and sarcomas.

49. The method of claim 1, wherein the cancer is selected from the group consisting of bone cancer, brain and CNS cancer, connective tissue cancer, esophageal cancer, eye cancer, Hodgkin's lymphoma, larynx cancer, oral cavity cancer, skin cancer, and testicular cancer.

50. The method of claim 47, further comprising administering an anti-cancer therapy.

51. The method of claim 50, wherein the anti-cancer therapy is an antibody.

52. The method of claim 47, wherein the subject is a human.

53. The method of claim 47, wherein the subject is selected from the group consisting of a dog, a cat, and a horse.

54. The method of claim 1, further comprising administering an antibody specific for a cell surface antigen, and wherein the immune response results in antigen dependent cellular cytotoxicity (ADCC).

55. The method of claim 1, wherein the subject has or is at risk of developing an infectious disease and wherein the method is a method for treating or preventing the infectious disease.

56. The method of claim 54, wherein the subject is a human.

57. The method of claim 54, further comprising administering an antigen to the subject.

58. The method of claim 57, wherein the antigen is selected from the group consisting of a bacterial antigen, a viral antigen, a parasitic antigen, and a fungal antigen.

59. The method of claim 56, wherein the subject is selected from the group consisting of a dog, cat, horse, cow, pig, sheep, goat, chicken, monkey, and fish.

60. The method of claim 59, further comprising administering an antigen to the subject.

61. The method of claim 59, wherein the antigen is derived from a microorganism selected from the group consisting of herpesviridae, retroviridae, orthomyxoviridae, toxoplasma, haemophilus, campylobacter, clostridium, E.coli, and staphylococcus.

5            62. The method of claim 1, wherein the immunostimulatory nucleic acid is a TG nucleic acid.

**63. The method of claim 62, wherein the TG nucleic acid comprises**



**64. The method of claim 62, wherein the TG nucleic acid comprises**



65. The method of claim 63, wherein N<sub>1</sub> is a nucleic acid sequence composed of a number of nucleotides ranging from (11-N<sub>2</sub>) to (21-N<sub>2</sub>).

66. The method of claim 63, wherein N<sub>2</sub> is a nucleic acid sequence composed of a number of nucleotides ranging from (11-N<sub>1</sub>) to (21-N<sub>1</sub>).

15 67. The method of claim 64, wherein N<sub>1</sub> is a nucleic acid sequence composed of a number of nucleotides ranging from (9-N<sub>2</sub>) to (19-N<sub>2</sub>).

68. The method of claim 64, wherein N<sub>2</sub> is a nucleic acid sequence composed of a number of nucleotides ranging from (9-N<sub>1</sub>) to (19-N<sub>1</sub>).

**69. The method of claim 63, wherein X<sub>2</sub> is thymidine.**

20 70. The method of claim 64, wherein X<sub>3</sub> is thymidine.

71. The method of claim 64, wherein  $X_1X_2$  are nucleotides selected from the group consisting of GT, GG, GA, AA, AT, AG, CT, CA, CG, TA and TT.

72. The method of claim 64, wherein X<sub>3</sub>X<sub>4</sub> are nucleotides selected from the group consisting of TT, CT, AT, AG, CG, TC, AC, CC, TA, AA, and CA.



- 163 -

73. The method of claim 63, wherein  $X_3X_4$  are nucleotides selected from the group consisting of TT, TC, TA and TG.

74. The method of claim 1, wherein the subject is at risk of developing cancer and the method is a method of preventing the cancer.

5        75. The method of claim 50, wherein the anti-cancer therapy is selected from the group consisting of a chemotherapeutic agent, an immunotherapeutic agent and a cancer vaccine.

76. A method for preventing disease in a subject, comprising:  
administering to the subject an immunostimulatory nucleic acid on a regular basis  
10 to prevent disease in the subject, wherein the immunostimulatory nucleic acid is selected from the group consisting of a T-rich nucleic acid and a TG nucleic acid.

77. A method for inducing an innate immune response, comprising  
administering to the subject an immunostimulatory nucleic acid in an amount  
effective for activating an innate immune response, wherein the immunostimulatory  
15 nucleic acid is selected from the group consisting of a T-rich nucleic acid and a TG nucleic acid.

78. A composition comprising  
a sustained release device including an immunostimulatory nucleic acid, wherein  
the immunostimulatory nucleic acid is free of unmethylated CpG motifs and is selected  
20 from the group consisting of a T-rich nucleic acid and a TG nucleic acid.

79. The composition of claim 78, wherein the immunostimulatory nucleic acid has a phosphodiester backbone.

80. A composition of a nutritional supplement comprising  
an immunostimulatory nucleic acid in a delivery device selected from the group  
25 consisting of a capsule, a pill, and a sublingual tablet, wherein the immunostimulatory nucleic acid is free of unmethylated CpG motifs and is selected from the group consisting of a T-rich nucleic acid and a TG nucleic acid.

- 164 -

81. The composition of claim 80, wherein the immunostimulatory nucleic acid has a phosphorothioate backbone.

82. A composition comprising  
an immunostimulatory nucleic acid and an antigen, wherein the  
5 immunostimulatory nucleic acid is free of unmethylated CpG motifs and is selected from the group consisting of a T-rich nucleic acid and a TG nucleic acid.

83. A composition comprising  
an immunostimulatory nucleic acid and an anti-microbial agent, wherein the  
immunostimulatory nucleic acid is free of unmethylated CpG motifs and is selected from  
10 the group consisting of a T-rich nucleic acid and a TG nucleic acid.

84. The composition of claim 83, wherein the anti-microbial agent is selected from the group consisting of an anti-viral agent, an anti-fungal agent, an anti-parasitic agent, and an anti-bacterial agent.

85. The method of claim 5, wherein the immunostimulatory nucleic acid  
15 comprises at least 3, at least 4, at least 5, at least 6, at least 7, or at least 8 T motifs.

86. The method of claim 5, wherein at least 2 of the plurality of poly T motifs each comprises at least three contiguous T nucleotide residues.

87. The method of claim 5, wherein at least two of the poly T motifs each comprises at least four contiguous T nucleotide residues.

20 88. The method of claim 5, wherein the plurality of poly T motifs is at least 3 motifs and wherein at least 3 motifs each comprises at least 3 contiguous T nucleotide residues.

89. The method of claim 5, wherein the plurality of poly T motifs is at least 4 motifs and wherein the at least 4 motifs each comprises at least 3 contiguous T  
25 nucleotide residues.

90. The method of claim 5, wherein at least one of the plurality of poly T motifs comprises at least 5, at least 6, at least 7, or at least 8 contiguous nucleotide residues.

- 165 -

91. The method of claim 1, wherein the immunostimulatory nucleic acid is free of two CpG dinucleotides.

92. The method of claim 1, wherein the immunostimulatory nucleic acid is free of three CpG dinucleotides.

5 93. The method of claim 1, wherein the immunostimulatory nucleic acid includes at least two poly C sequences of at least 3 contiguous C nucleotide residues.

94. The method of claim 1, wherein the immunostimulatory nucleic acid is free of two poly A sequences of at least 3 contiguous A nucleotide residues.

95. A pharmaceutical composition comprising an effective amount for  
10 stimulating an immune response of an isolated immunostimulatory nucleic acid of claim 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 32, 40, 41, 64, 65, 66, 67, 68, 69, 70, 71, 72, 85, 86, 87, 88, 89, 90, 91, 92, 93, or 94 and a pharmaceutically acceptable carrier.

96. A composition of matter, comprising an isolated immunostimulatory nucleic  
15 acid of claim 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 32, 40, 41, 64, 65, 66, 67, 68, 69, 70, 71, 72, 85, 86, 87, 88, 89, 90, 91, 92, 93, or 94 and a pharmaceutically acceptable carrier.

97 The method of claim 80 wherein the nucleic acid further comprises a plurality  
of CpG motifs, and wherein the plurality is at least 3 motifs, at least 4 motifs and  
20 wherein the at least 4 motifs each comprises at least 3 contiguous T nucleotide residues.

98 The method of claim 90 wherein the plurality of CpG motifs and poly T motifs are interspersed.

99. A composition, comprising:  
an immunostimulatory nucleic acid and an anti-cancer therapy,  
25 formulated in a pharmaceutically-acceptable carrier and in an effective amount to treat a cancer or to reduce the risk of developing a cancer, wherein the immunostimulatory nucleic acid is selected from the group consisting of a T-rich nucleic acid and a TG nucleic acid.

100. A composition, comprising:

- 166 -

an immunostimulatory nucleic acid and an asthma/allergy medicament, formulated in a pharmaceutically-acceptable carrier and in an effective amount for preventing or treating an immune response associated with exposure to a mediator of asthma or allergy, wherein the immunostimulatory nucleic acid is selected from the group consisting of a T-rich nucleic acid, a TG nucleic acid and a C-rich nucleic acid.

101. A composition comprising

an immunostimulatory nucleic acid selected from the group consisting of SEQ ID NO: 95-136, SEQ ID NO: 138-152, SEQ ID NO: 154-222, SEQ ID NO: 224-245, SEQ ID NO: 247-261, SEQ ID NO: 263-299, SEQ ID NO: 301, SEQ ID NO: 303-4109, SEQ ID NO: 414-420, SEQ ID NO: 424, SEQ ID NO: 426-947, SEQ ID NO: 959-1022, SEQ ID NO: 1024-1093, and a pharmaceutically acceptable carrier.

102. A composition comprising

an immunostimulatory nucleic acid consisting essentially of:

5' M<sub>1</sub>TCGTCGTTM<sub>2</sub> 3'

wherein at least one of the Cs is unmethylated, wherein M<sub>1</sub> is a nucleic acid having at least one nucleotide, wherein M<sub>2</sub> is a nucleic acid having between 0 and 50 nucleotides, and wherein the immunostimulatory nucleic acid has less than 100 nucleotides.

103. A pharmaceutical composition comprising

an immunostimulatory nucleic acid comprising:

5' TCGTCGTT 3'

wherein at least one of the Cs is unmethylated, wherein the immunostimulatory nucleic acid has less than 100 nucleotides and a phosphodiester backbone, and

a sustained release device.

104. The pharmaceutical composition of claim 103 wherein the sustained release device is a microparticle.

105. The pharmaceutical composition of claim 103, further comprising an antigen.

1/10

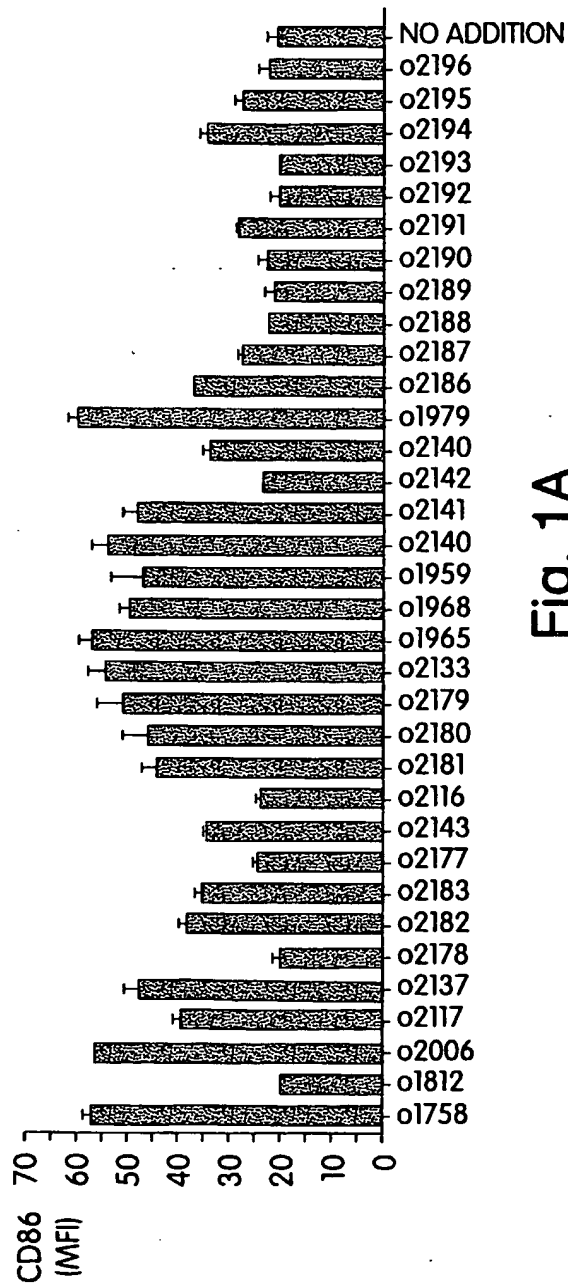


Fig. 1A

2/10

## STATISTICAL ANALYSIS

	MEAN	STD.DEV.	STD.ERROR	NUMBER	MINIMUM	MAXIMUM	# MISSING
o1758	57.60	.71	.50	2	57.10	58.10	2
o1812	20.70	.57	.40	2	20.30	21.10	2
o2006	56.35	.35	.25	2	56.10	56.60	2
o2117	40.00	1.13	.80	2	39.20	40.80	2
o2137	47.60	3.39	2.40	2	45.20	50.00	2
o2178	21.00	.99	.70	2	20.30	21.70	2
o2182	37.75	.35	.25	2	37.50	38.00	2
o2183	35.30	1.84	1.30	2	34.00	36.60	2
o2177	25.00	.71	.50	2	24.50	25.50	2
o2143	34.70	.71	.50	2	34.20	35.20	2
o2116	24.35	.64	.45	2	23.90	24.80	2
o2181	44.25	3.18	2.25	2	42.00	46.50	2
o2180	45.90	5.94	4.20	2	41.70	50.10	2
o2179	50.70	6.93	4.90	2	45.80	55.60	2
o2133	53.75	4.31	3.05	2	50.70	56.80	2
o1965	56.20	3.82	2.70	2	53.50	58.90	2
o1968	49.35	1.91	1.35	2	48.00	50.70	2
o2159	46.80	7.92	5.60	2	41.20	52.40	2
o2140	53.25	4.74	3.35	2	49.90	56.60	2
o2141	47.40	4.10	2.90	2	44.50	50.30	2
o2142	23.20	.42	.30	2	22.90	23.50	2
o1840	33.50	1.13	.80	2	32.70	34.30	2
o1979	59.50	1.70	1.20	2	58.30	60.70	2
o2186	36.90	.14	.10	2	36.80	37.00	2
o2187	27.15	.78	.55	2	26.60	27.70	2
o2188	22.25	.21	.15	2	22.10	22.40	2
o2189	21.45	1.20	.85	2	20.60	22.30	2
o2190	22.95	1.20	.85	2	22.10	23.80	2
o2191	28.35	.49	.35	2	28.00	28.70	2
o2192	20.40	1.70	1.20	2	19.20	21.60	2
o2193	19.70	.57	.40	2	19.30	20.10	2
o2194	34.00	1.70	1.20	2	32.80	35.20	2
o2195	27.30	1.27	.90	2	26.40	28.20	2
o2196	22.45	2.76	1.95	2	20.50	24.40	2
NO ADDITION	20.90	2.77	1.38	4	18.60	24.90	0

Fig. 1B

3/10

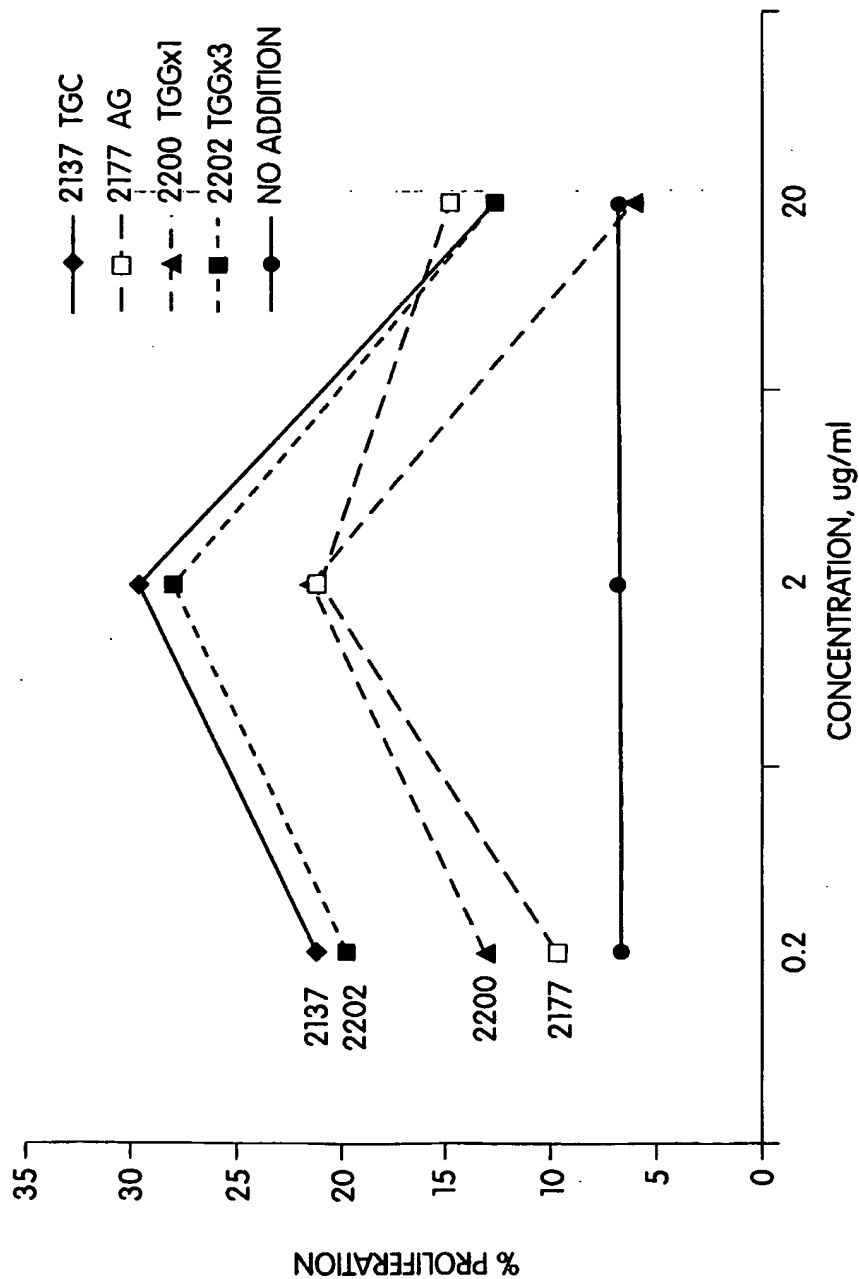


Fig. 2

4/10

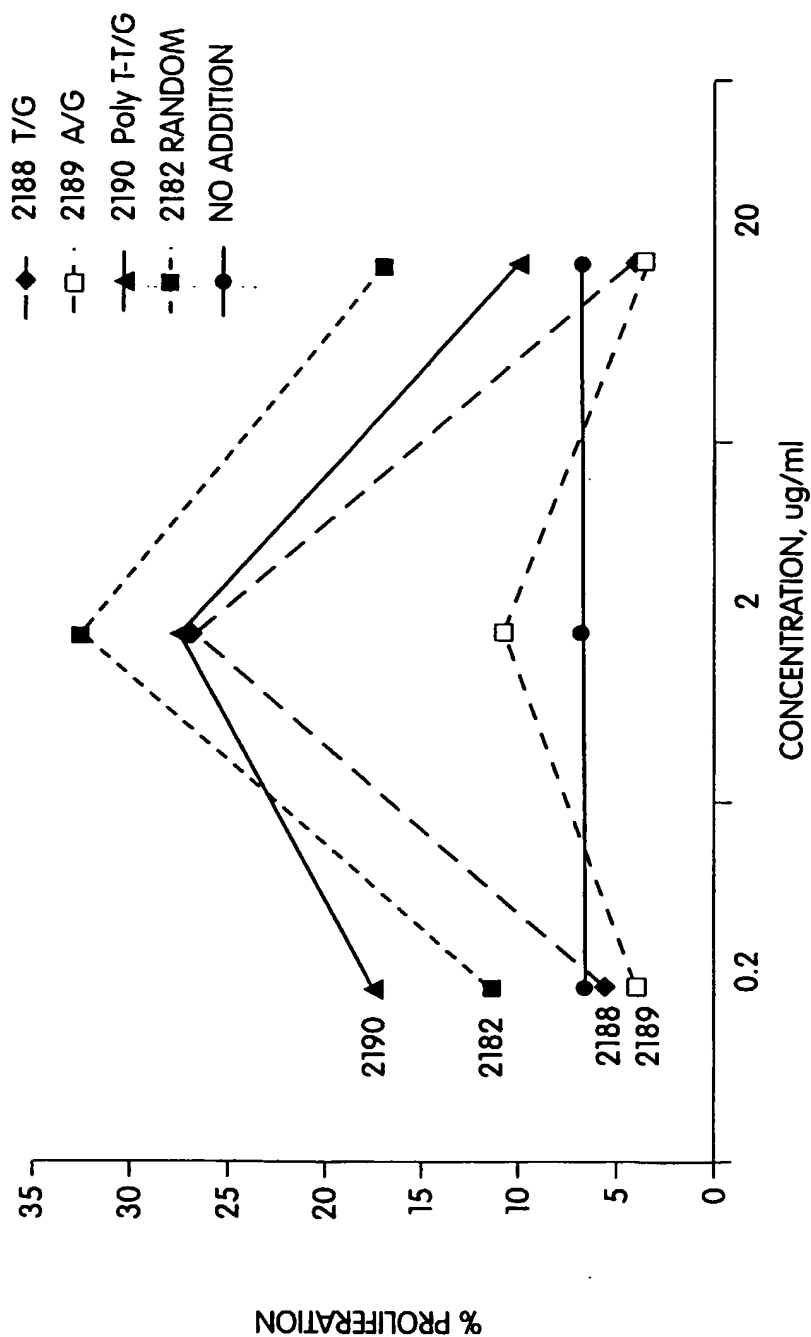


Fig. 3



5/10

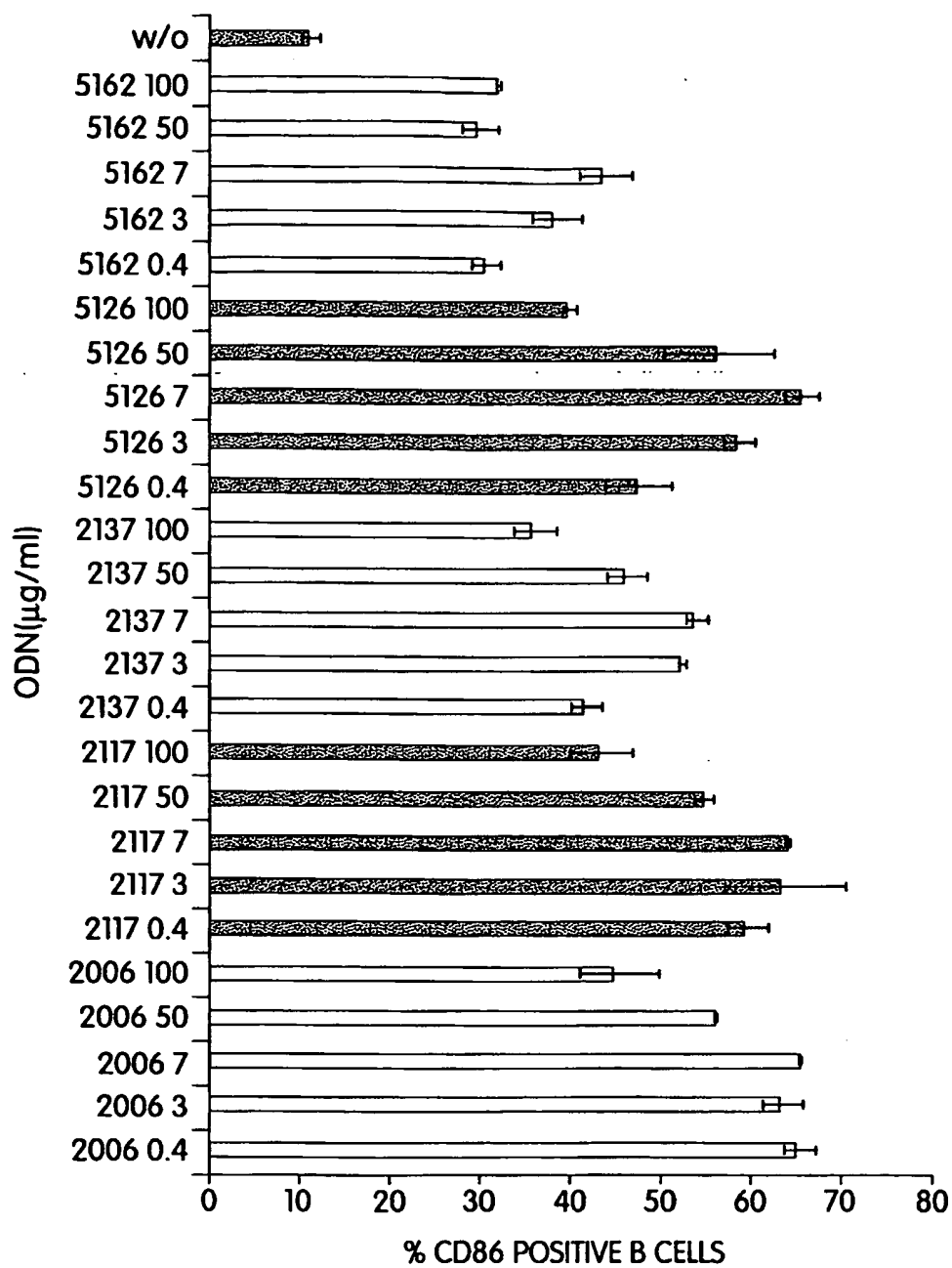


Fig. 4

6/10

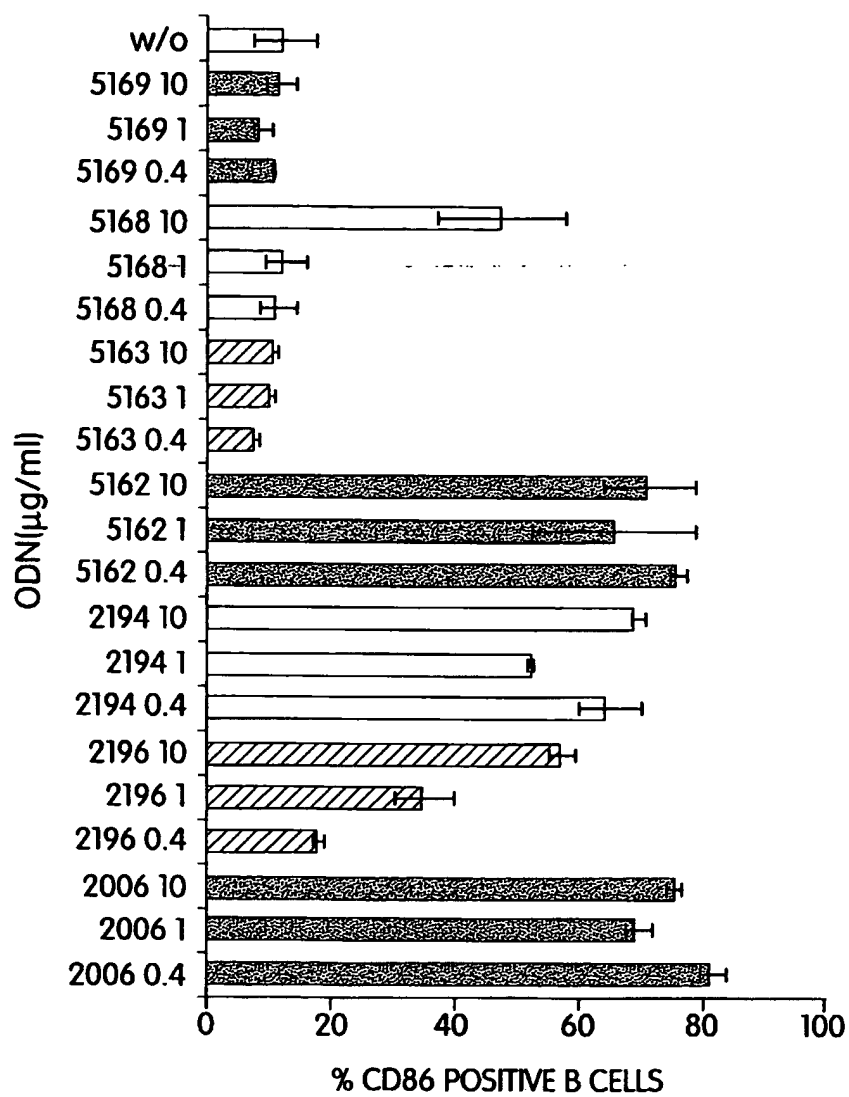


Fig. 5

7/10

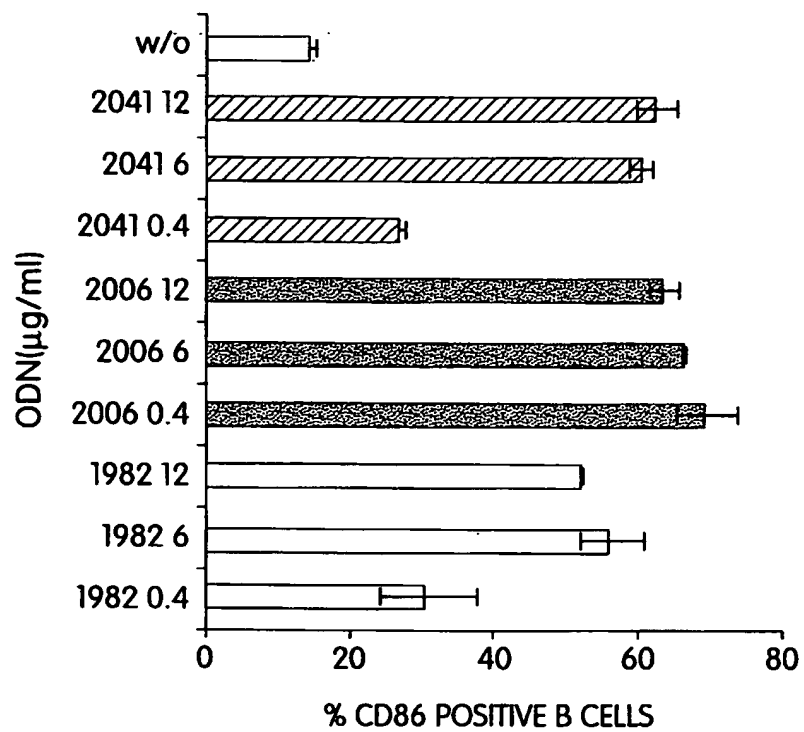


Fig. 6

8/10

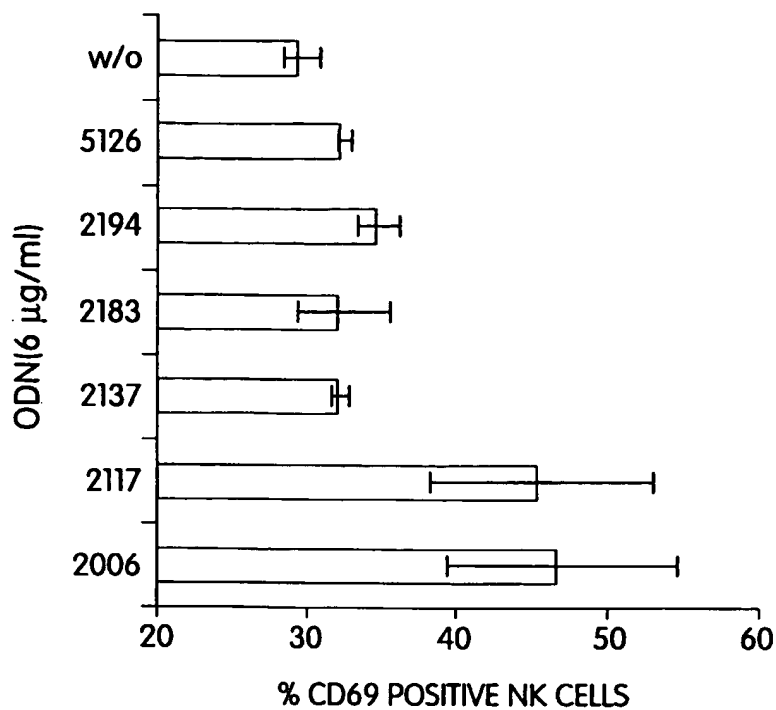


Fig. 7

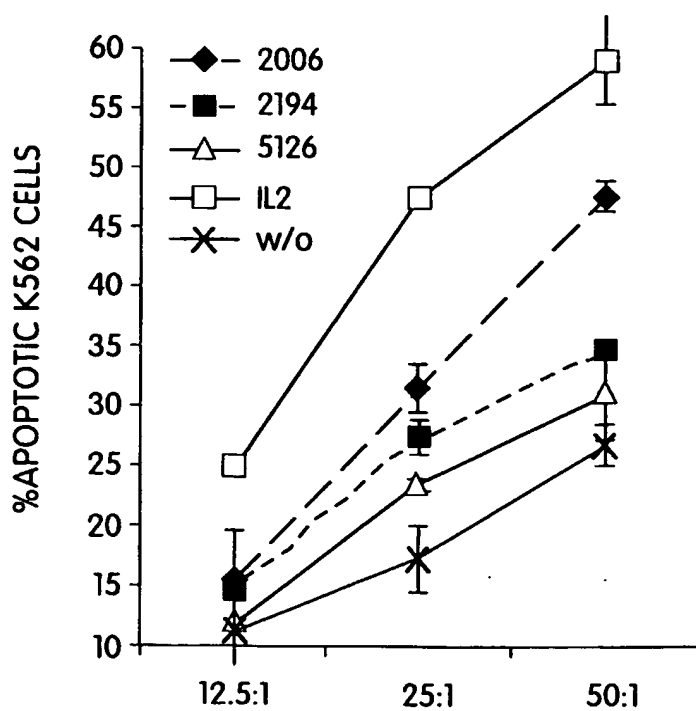


Fig. 8

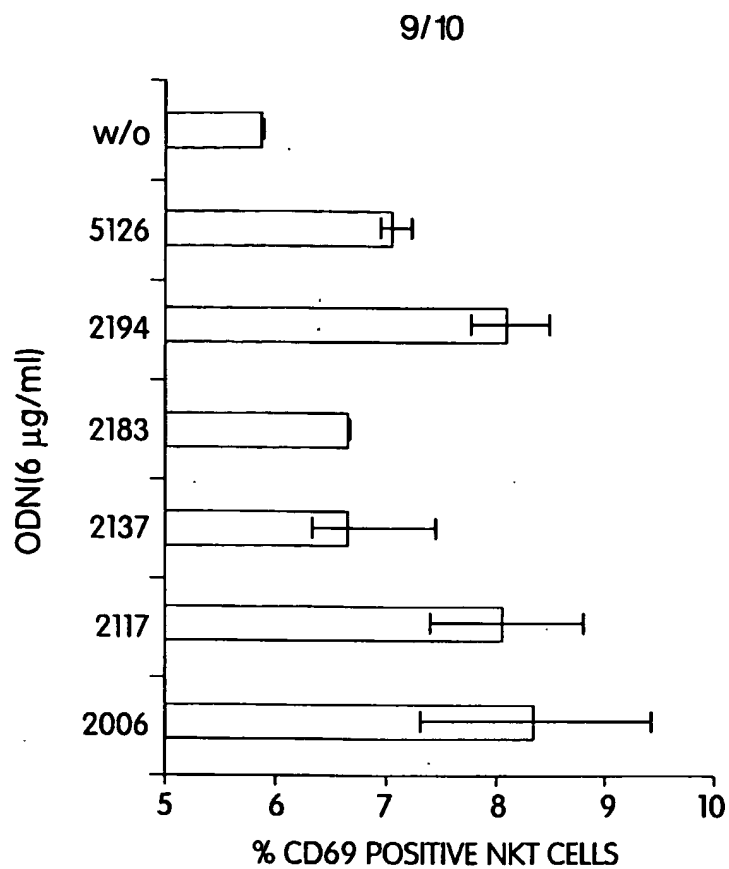


Fig. 9

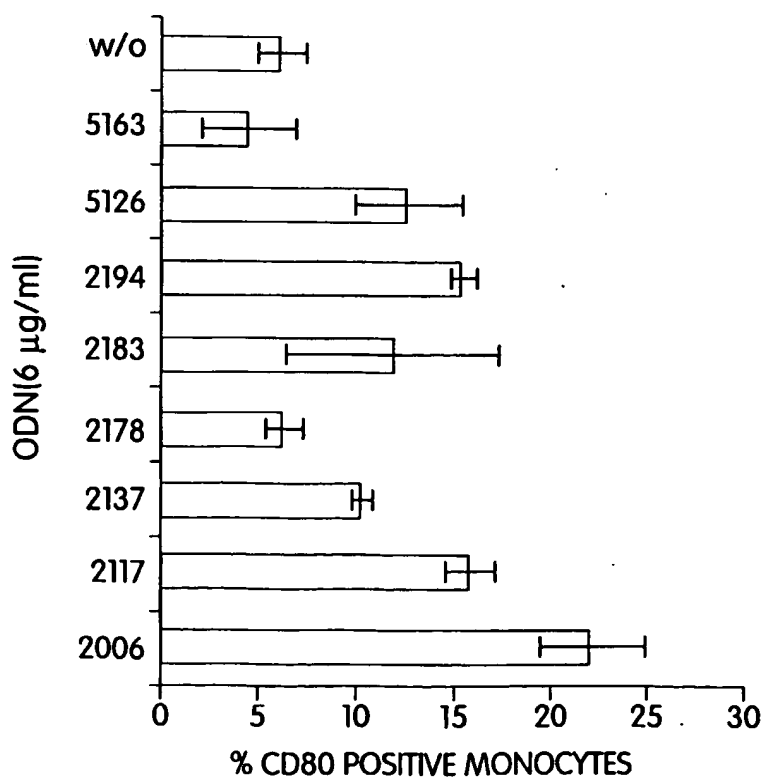


Fig. 10

10/10

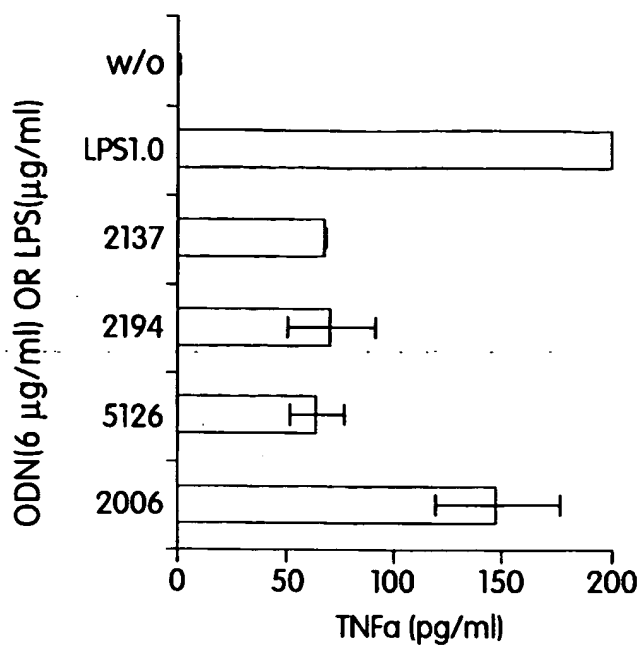


Fig. 11

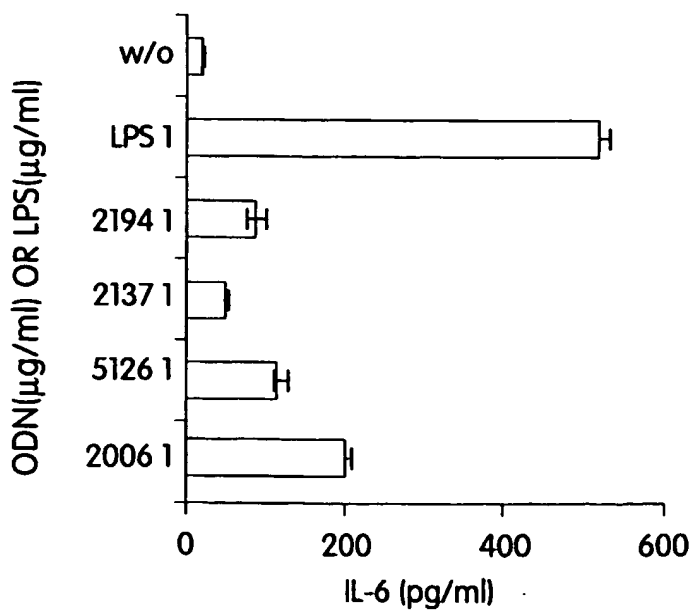


Fig. 12

## SEQUENCE LISTING

<110> University of Iowa Research Foundation  
Coley Pharmaceutical Group GmbH

<120> Immunostimulatory Nucleic Acids

<130> C1039/7035WO (HCL/MAT)

<150> US 60/156,113  
<151> 1999-09-25

<150> US 60/156,135  
<151> 1999-09-27

<150> US 60/227,436  
<151> 2000-08-23

<160> 1145

<170> FastSEQ for Windows Version 3.0

<210> 1  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Sequence

<400> 1  
tctcccagcg tgcgccat 18

<210> 2  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 2  
ataatccagc ttgaaccaag 20

<210> 3  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 3  
ataatcgacg ttcaagcaag 20

<210> 4  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 4  
taccgctgc gaccctct 18

<210> 5  
<211> 9  
<212> DNA

<213> Artificial Sequence  
 <400> 5  
 ggggaggggt 9  
 <210> 6  
 <211> 9  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 6  
 ggggagggg 9  
 <210> 7  
 <211> 9  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 7  
 ggtgaggtg 9  
 <210> 8  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <220>  
 <221> modified\_base  
 <222> (8)...(8)  
 <223> m5c  
 <400> 8  
 tccatgtngt tcctgatgct 20  
 <210> 9  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence  
 <220>  
 <221> modified\_base  
 <222> (11)...(11)  
 <223> m5c  
 <400> 9  
 gctaccttag ngtaga 15  
 <210> 10  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <220>  
 <221> modified\_base  
 <222> (8)...(8)  
 <223> m5c  
 <400> 10  
 tccatgangt tcctgatgct 20  
 <210> 11



<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (13)...(13)  
<223> m5c

<400> 11  
tccatgacgt tcntgatgct 20

<210> 12  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (7)...(7)  
<223> m5c

<400> 12  
gctagangtt agtgt 15

<210> 13  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 13  
agctccatgg tgctcactg 19

<210> 14  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 14  
ccacgtcgac cctcaggcga 20

<210> 15  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 15  
gcacatcgtc ccgcagccga 20

<210> 16  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 16  
gtcactcgtg gtacctcga 19

<210> 17  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 17  
gttggataca ggccagactt tgttg 25

<210> 18  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 18  
gattcaactt gcgctcatct taggc 25

<210> 19  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 19  
accatggacg aactgtttcc cctc 24

<210> 20  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 20  
accatggacg agctgtttcc cctc 24

<210> 21  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 21  
accatggacg acctgtttcc cctc 24

<210> 22  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 22  
accatggacg tactgtttcc cctc 24

<210> 23  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 23  
accatggacg gtctgtttcc cctc 24

<210> 24  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 24  
accatggacg ttctgtttcc cctc 24

<210> 25

<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 25  
ccactcacat ctgctgctcc acaag 25

<210> 26  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 26  
acttctcata gtccctttgg tccag 25

<210> 27  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 27  
tccatgagct tcctgagtct 20

<210> 28  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (9)...(9)  
<223> I

<221> modified\_base  
<222> (11)...(11)  
<223> I

<221> modified\_base  
<222> (15)...(15)  
<223> I

<400> 28  
gaggaaggng nggangacgt 20

<210> 29  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (7)...(7)  
<223> I

<221> modified\_base  
<222> (13)...(13)  
<223> I

<221> modified\_base  
<222> (18)...(18)  
<223> I

<400> 29	
gtgaatncgt tcncgggnct	20
<210> 30	
<211> 6	
<212> DNA	
<213> Artificial Sequence	
<400> 30	
aaaaaa	6
<210> 31	
<211> 6	
<212> DNA	
<213> Artificial Sequence	
<400> 31	
cccccc	6
<210> 32	
<211> 6	
<212> DNA	
<213> Artificial Sequence	
<400> 32	
ctgtca	6
<210> 33	
<211> 6	
<212> DNA	
<213> Artificial Sequence	
<400> 33	
tcgtag	6
<210> 34	
<211> 6	
<212> DNA	
<213> Artificial Sequence	
<400> 34	
tcgtgg	6
<210> 35	
<211> 6	
<212> DNA	
<213> Artificial Sequence	
<400> 35	
cgtcgt	6
<210> 36	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<400> 36	
tccatgtcgg tcctgagtct	20
<210> 37	

<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 37  
tccatgccgg tcctgagtct 20

<210> 38  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 38  
tccatgacgg tcctgagtct 20

<210> 39  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 39  
tccatgacgg tcctgagtct 20

<210> 40  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 40  
tccatgtcga tcctgagtct 20

<210> 41  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 41  
tccatgtcgc tcctgagtct 20

<210> 42  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 42  
tccatgtcgt tcctgagtct 20

<210> 43  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 43  
tccatgacgt tcctgagtct 20

<210> 44  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 44

8

tccataacgt tcctgagtct

20

&lt;210&gt; 45

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 45

tccatgacgt ccctgagtct

20

&lt;210&gt; 46

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 46

tccatcacgt gcctgagtct

20

&lt;210&gt; 47

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 47

tccatgctgg tcctgagtct

20

&lt;210&gt; 48

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;221&gt; modified\_base

&lt;222&gt; (8)...(8)

&lt;223&gt; m5c

&lt;400&gt; 48

tccatgtngg tcctgagtct

20

&lt;210&gt; 49

&lt;211&gt; 39

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 49

ccgcttcctc cagatgagct catgggtttc tccaccaag

39

&lt;210&gt; 50

&lt;211&gt; 39

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 50

cttggtggag aaacccatga gctcatctgg aggaagcgg

39

&lt;210&gt; 51

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 51

ccccaaggg atgagaagtt

20

&lt;210&gt; 52

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 52

agatagcaaa tcggctgacg

20

&lt;210&gt; 53

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 53

ggttcacgtg ctcatggctg

20

&lt;210&gt; 54

&lt;211&gt; 18

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 54

tctcccagcg tgcgccat

18

&lt;210&gt; 55

&lt;211&gt; 18

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 55

tctcccagcg tgcgccat

18

&lt;210&gt; 56

&lt;211&gt; 18

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 56

taccggtgc gaccctct

18

&lt;210&gt; 57

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 57

ataatccagc ttgaaccaag

20

&lt;210&gt; 58

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 58

ataatcgacg ttcaagcaag

20

&lt;210&gt; 59

&lt;211&gt; 20

&lt;212&gt; DNA

<213> Artificial Sequence

<400> 59  
tccatgattt tcctgatttt 20

<210> 60  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 60  
ttgttttttt gtttttttgt tttt 24

<210> 61  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 61  
ttttttttgt ttttttgttt tt 22

<210> 62  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 62  
tgctgctttt gtgcttttgt gctt 24

<210> 63  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 63  
tgctgcttgt gcttttgtgc tt 22

<210> 64  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 64  
gcattcatca ggcgggcaag aat 23

<210> 65  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 65  
taccgagctt cgacgagatt tca 23

<210> 66  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 66  
gcatgacgtt gagct 15



<210> 67  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 67  
cacgttgagg ggcac 15

<210> 68  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 68  
ctgctgagac tggag 15

<210> 69  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 69  
tccatgacgt tcctgacgtt 20

<210> 70  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 70  
gcatgagctt gagctga 17

<210> 71  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 71  
tcagcgtgcg cc 12

<210> 72  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 72  
atgacgttcc tgacgtt 17

<210> 73  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 73  
ttttgggggtt ttgggggttt 20

<210> 74  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 74  
tctaggcttt ttaggcttcc 20

<210> 75  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 75  
tgcatttttt aggccaccat 20

<210> 76  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 76  
tctcccagcg tgcgtgcgcc at 22

<210> 77  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 77  
tctcccagcg ggcgcat 17

<210> 78  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 78  
tctcccagcg agcgccat 18

<210> 79  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 79  
tctcccagcg cgcgccat 18

<210> 80  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 80  
ggggtgacgt tcagggggg 19

<210> 81  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 81  
ggggtccagc gtgcgccatg gggg 24

<210> 82  
<211> 19

<212> DNA  
 <213> Artificial Sequence

<400> 82  
 ggggtgtcgt tcagggggg 19

<210> 83  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<400> 83  
 tccatgtcgt tcctgtcgtt 20

<210> 84  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<400> 84  
 tccatagcgt tcctagcgtt 20

<210> 85  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<400> 85  
 tcgtcgctgt ctccgttct t 21

<210> 86  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence

<400> 86  
 gcatgacgtt gagct 15

<210> 87  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<400> 87  
 tctcccagcg tgcgcatat 20

<210> 88  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> modified\_base  
 <222> (8)...(8)  
 <223> m5c

<221> modified\_base  
 <222> (17)...(17)  
 <223> m5c

<400> 88

		14
tccatgangt tcctgangtt		20
<210> 89		
<211> 15		
<212> DNA		
<213> Artificial Sequence		
<220>		
<221> modified_base		
<222> (7)...(7)		
<223> m5c		
<400> 89		
gcatgangtt gagct		15
<210> 90		
<211> 16		
<212> DNA		
<213> Artificial Sequence		
<400> 90		
tccagcgtgc gccata		16
<210> 91		
<211> 18		
<212> DNA		
<213> Artificial Sequence		
<400> 91		
tctcccagcg tgcgccat		18
<210> 92		
<211> 20		
<212> DNA		
<213> Artificial Sequence		
<400> 92		
tccatgagct tcctgagtct		20
<210> 93		
<211> 15		
<212> DNA		
<213> Artificial Sequence		
<400> 93		
gcatgtcggt gagct		15
<210> 94		
<211> 19		
<212> DNA		
<213> Artificial Sequence		
<400> 94		
tcctgacgtt cctgacgtt		19
<210> 95		
<211> 15		
<212> DNA		
<213> Artificial Sequence		
<400> 95		

15

gcatgatggt gagct

15

&lt;210&gt; 96

&lt;211&gt; 15

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 96

gcatttcgag gagct

15

&lt;210&gt; 97

&lt;211&gt; 15

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 97

gcatgtagct gagct

15

&lt;210&gt; 98

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 98

tccaggacgt tcctagttct

20

&lt;210&gt; 99

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 99

tccaggagct tcctagttct

20

&lt;210&gt; 100

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 100

tccaggatgt tcctagttct

20

&lt;210&gt; 101

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 101

tccagtctag gcctagttct

20

&lt;210&gt; 102

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 102

tccagttcga gcctagttct

20

&lt;210&gt; 103

&lt;211&gt; 15

&lt;212&gt; DNA

<213> Artificial Sequence

<400> 103  
gcatggcggtt gagct 15

<210> 104  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 104  
gcatagcggtt gagct 15

<210> 105  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 105  
gcattgcggtt gagct 15

<210> 106  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 106  
gcttgcgttg cgttt 15

<210> 107  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 107  
tctcccagcg ttgcgccata t 21

<210> 108  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 108  
tctcccagcg tgcgttatat 20

<210> 109  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 109  
tctccctgcg tgcgccatat 20

<210> 110  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 110  
tctgctgctg tgcgccatat 20

<210> 111  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 111  
tctcctagcg tgcgccatat 20

<210> 112  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 112  
tctcccagcg tgcgcctttt 20

<210> 113  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_difference  
<222> (5)...(5)  
<223> n is a or g or c or t/u

<221> misc\_difference  
<222> (6)...(6)  
<223> d is a or g or t/u; not c

<221> misc\_difference  
<222> (9)...(10)  
<223> h is a or c or t/u; not g

<400> 113  
gctandcghh agc 13

<210> 114  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 114  
tcctgacgtt ccc 13

<210> 115  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 115  
ggaagacgtt aga 13

<210> 116  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 116  
tcctgacgtt aga 13

<210> 117  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 117  
tcagaccagc tggtcgggtg ttcctga 27

<210> 118  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 118  
tcaggaacac ccgaccagct ggtctga 27

<210> 119  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 119  
gctagtcgat agc 13

<210> 120  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 120  
gctagtcgct agc 13

<210> 121  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 121  
gcttgacgtc tagc 14

<210> 122  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 122  
gcttgacggt tagc 14

<210> 123  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 123  
gcttgacgtc aagc 14

<210> 124  
<211> 14  
<212> DNA  
<213> Artificial Sequence



<400> 124  
gctagacggt tagc 14

<210> 125  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 125  
tccatgacat tcctgatgct 20

<210> 126  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 126  
gctagacgtc tagc 14

<210> 127  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 127  
ggctatgtcg ttcctagcc 19

<210> 128  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 128  
ggctatgtcg atcctagcc 19

<210> 129  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 129  
ctcatgggtt tctccaccaa g 21

<210> 130  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 130  
cttggtggag aaacccatga g 21

<210> 131  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 131  
tccatgacgt tcctagttct 20

<210> 132  
<211> 24

<212> DNA  
<213> Artificial Sequence

<400> 132  
ccgcttcctc cagatgagct catg 24

<210> 133  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 133  
catgagctca tctggaggaa gcgg 24

<210> 134  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 134  
ccagatgagc tcatgggttt ctcc 24

<210> 135  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 135  
ggagaaacct atgagctcat ctgg 24

<210> 136  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 136  
agcatcagga acgacatgga 20

<210> 137  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 137  
tccatgacgt tcctgacgtt 20

<210> 138  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 138  
gcgcgcgcgc gcgcgcgcg 19

<210> 139  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 139  
ccggccggcc ggccggccgg 20

<210> 140  
<211> 43  
<212> DNA  
<213> Artificial Sequence

<400> 140  
ttccaatcag cccacccgc tctggccca cctcaccct cca 43

<210> 141  
<211> 43  
<212> DNA  
<213> Artificial Sequence

<400> 141  
tggagggtga gggtagggcc agagcgggtg gggctgattg gaa 43

<210> 142  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 142  
tcaaattgtg gattttcca tgagtct 27

<210> 143  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 143  
agactcatgg gaaaatcca catttga 27

<210> 144  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 144  
tgccaagtgc tgagtcacta ataaaga 27

<210> 145  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 145  
tctttattag tgactcagca cttggca 27

<210> 146  
<211> 31  
<212> DNA  
<213> Artificial Sequence

<400> 146  
tgcaggagt ccgggttttc cccaaccccc c 31

<210> 147  
<211> 31  
<212> DNA  
<213> Artificial Sequence

<400> 147  
ggggggttg ggaaaacccg gacttcctgc a 31

<210> 148  
<211> 38  
<212> DNA  
<213> Artificial Sequence

<400> 148  
ggggactttc cgctggggac tttccagggg gactttcc 38

<210> 149  
<211> 45  
<212> DNA  
<213> Artificial Sequence

<400> 149  
tccatgacgt tccttccat gacgttcctc tccatgacgt tcctc 45

<210> 150  
<211> 45  
<212> DNA  
<213> Artificial Sequence

<400> 150  
gaggaacgtc atggagagga acgtcatgga gaggaacgtc atgga 45

<210> 151  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 151  
ataatagagc ttcaagcaag 20

<210> 152  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 152  
tccatgacgt tcctgacgtt 20

<210> 153  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 153  
tccatgacgt tcctgacgtt 20

<210> 154  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 154  
tccaggactt tcctcaggtt 20

<210> 155

<211> 45  
<212> DNA  
<213> Artificial Sequence

<400> 155  
tcttgcgatg ctaaaggacg tcacattgca caatcttaat aaggt 45

<210> 156  
<211> 45  
<212> DNA  
<213> Artificial Sequence

<400> 156  
accttattaa gattgtgcaa tgtgacgtcc tttagcatcg caaga 45

<210> 157  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<400> 157  
tcctgacgtt cctggcggtc ctgtcgct 28

<210> 158  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 158  
tcctgtcgct cctgtcgct 19

<210> 159  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 159  
tcctgacgtt gaagt 15

<210> 160  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 160  
tcctgtcggt gaagt 15

<210> 161  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 161  
tcctggcggt gaagt 15

<210> 162  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 162

tcctgccgtt gaagt	15
<210> 163	
<211> 15	
<212> DNA	
<213> Artificial Sequence	
<400> 163	
tccttacgtt gaagt	15
<210> 164	
<211> 15	
<212> DNA	
<213> Artificial Sequence	
<400> 164	
tcctaacgtt gaagt	15
<210> 165	
<211> 15	
<212> DNA	
<213> Artificial Sequence	
<400> 165	
tcctcacgtt gaagt	15
<210> 166	
<211> 15	
<212> DNA	
<213> Artificial Sequence	
<400> 166	
tcctgacgat gaagt	15
<210> 167	
<211> 15	
<212> DNA	
<213> Artificial Sequence	
<400> 167	
tcctgacgct gaagt	15
<210> 168	
<211> 15	
<212> DNA	
<213> Artificial Sequence	
<400> 168	
tcctgacggt gaagt	15
<210> 169	
<211> 15	
<212> DNA	
<213> Artificial Sequence	
<400> 169	
tcctgacgta gaagt	15
<210> 170	
<211> 15	
<212> DNA	

<213> Artificial Sequence  
 <400> 170  
 tcctgacgtc gaagt 15  
 <210> 171  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 171  
 tcctgacgtg gaagt 15  
 <210> 172  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 172  
 tcctgagctt gaagt 15  
 <210> 173  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 173  
 gggggacgtt ggggg 15  
 <210> 174  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 174  
 tcctgacgtt ccttc 15  
 <210> 175  
 <211> 22  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 175  
 tctcccagcg agcgagcgcc at 22  
 <210> 176  
 <211> 32  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 176  
 tcctgacgtt ccctggcgg tccctgtcg ct 32  
 <210> 177  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 177  
 tcctgtcgct cctgtcgctc ctgtcgct 28

<210> 178  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 178  
tcctggcggg gaagt 15

<210> 179  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (7)...(7)  
<223> m5c

<400> 179  
tcctgangtt gaagt 15

<210> 180  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (3)...(3)  
<223> m5c

<400> 180  
tcntgacgtt gaagt 15

<210> 181  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 181  
tcctagcgtt gaagt 15

<210> 182  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 182  
tccagacgtt gaagt 15

<210> 183  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 183  
tcctgacggg gaagt 15

<210> 184  
<211> 15  
<212> DNA



<213> Artificial Sequence	
<400> 184	
tcctggcggg gaagt	15
<210> 185	
<211> 27	
<212> DNA	
<213> Artificial Sequence	
<400> 185	
ggctccgggg agggaaatddd tgtctat	27
<210> 186	
<211> 27	
<212> DNA	
<213> Artificial Sequence	
<400> 186	
atagacaaaa attccctccc cggagcc	27
<210> 187	
<211> 21	
<212> DNA	
<213> Artificial Sequence	
<400> 187	
tccatgagct tccttgagtc t	21
<210> 188	
<211> 21	
<212> DNA	
<213> Artificial Sequence	
<400> 188	
tcgtcgctgt ctccgcttct t	21
<210> 189	
<211> 21	
<212> DNA	
<213> Artificial Sequence	
<400> 189	
tcgtcgctgt ctccgcttct t	21
<210> 190	
<211> 23	
<212> DNA	
<213> Artificial Sequence	
<400> 190	
tcgagacatt gcacaatcat ctg	23
<210> 191	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<400> 191	
cagattgtgc aatgtctcga	20

<210> 192  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 192  
tccatgtcgt tcctgatgcg 20

<210> 193  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 193  
gcgatgtcgt tcctgatgct 20

<210> 194  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 194  
gcgatgtcgt tcctgatgcg 20

<210> 195  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 195  
tccatgtcgt tccgcgcgcg 20

<210> 196  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 196  
tccatgtcgt tcctgccgct 20

<210> 197  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 197  
tccatgtcgt tcctgtagct 20

<210> 198  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 198  
gcggcgggcg gcgcgcgccc 20

<210> 199  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 199  
atcaggaacg tcatgggaag c 21

<210> 200  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 200  
tccatgagct tcctgagtct 20

<210> 201  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 201  
tcaacggtt 8

<210> 202  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 202  
tcaagctt 8

<210> 203  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 203  
tcctgtcggt cctgtcggt 19

<210> 204  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 204  
tccatgtcgt tttgtcggt 20

<210> 205  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 205  
tcctgtcggt cctgtcggt 20

<210> 206  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 206  
tcctgtcgt tcctgtcggt 20

<210> 207  
<211> 29

<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 207  
tccattccat gacgttcctg atgcttcca 29

<210> 208  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 208  
tcctgtcggt tttgtggtt 20

<210> 209  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 209  
tcgtcgctgt ctccgttct t 21

<210> 210  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 210  
tcgtcgctgt ctgcccttct t 21

<210> 211  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 211  
tcgtcgctgt tgtcgtttct t 21

<210> 212  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<400> 212  
tcctgtcggt cctgtcggtg gaacgacagg 30

<210> 213  
<211> 40  
<212> DNA  
<213> Artificial Sequence

<400> 213  
tcctgtcggt cctgtcggtt caacgtcagg aacgacagga 40

<210> 214  
<211> 21

<212> DNA  
<213> Artificial Sequence

<400> 214  
ggggtctgtc gttttgggg g 21

<210> 215  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 215  
ggggtctgtg cttttgggg g 21

<210> 216  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 216  
tccggccggt gaagt 15

<210> 217  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 217  
tccggacggt gaagt 15

<210> 218  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 218  
tcccgccggt gaagt 15

<210> 219  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 219  
tccagacggt gaagt 15

<210> 220  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 220  
tcccgacggt gaagt 15

<210> 221  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 221  
tccagagctt gaagt 15

<210> 222  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (8)...(8)  
<223> m5c

<221> modified\_base  
<222> (17)...(17)  
<223> m5c

<400> 222  
tccatgtngt tcctgtngtt 20

<210> 223  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 223  
tccatgacgt tcctgacgtt 20

<210> 224  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 224  
ggggttgacg ttttggggg 20

<210> 225  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 225  
tccaggactt ctctcaggtt 20

<210> 226  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 226  
tttttttttt tttttttttt 20

<210> 227  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 227  
tccatgccgt tcctgccgtt 20

<210> 228  
<211> 20  
<212> DNA

<213> Artificial Sequence

<400> 228  
tccatggcgg gcctggcggg 20

<210> 229  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 229  
tccatgacgt tcctgccgtt 20

<210> 230  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 230  
tccatgacgt tcctggcggg 20

<210> 231  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 231  
tccatgacgt tcctgcgttt 20

<210> 232  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 232  
tccatgacgg tcctgacggt 20

<210> 233  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 233  
tccatgcgtg cgtgcgtttt 20

<210> 234  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 234  
tccatgcgtt gcgttgcgtt 20

<210> 235  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)

<223> Biotin moiety attached at 5' end of sequence.

<400> 235  
tccattccat tctaggcctg agtcttccat 30

<210> 236  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 236  
tccatagcgt tcctagcggt 20

<210> 237  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 237  
tccatgtcgt tcctgtcggt 20

<210> 238  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 238  
tccatagcga tcctagcgat 20

<210> 239  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 239  
tccattgcgt tccttgcgtt 20

<210> 240  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 240  
tccatagcgg tcctagcggt 20

<210> 241  
<211> 29  
<212> DNA  
<213> Artificial Sequence

<400> 241  
tccatgattt tcctgcagtt cctgatttt 29

<210> 242  
<211> 29  
<212> DNA  
<213> Artificial Sequence

<400> 242  
tccatgacgt tcctgcagtt cctgacgtt 29



<210> 243  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 243  
ggcggcggcg gcggcggcgg 20

<210> 244  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 244  
tccacgacgt ttccgacgtt 20

<210> 245  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 245  
tcgtcgttgt cgtgtcgtt 20

<210> 246  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 246  
tcgtcgtttt gtcgttttgt cgtt 24

<210> 247  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 247  
tcgtcgttgt cgttttgtcg tt 22

<210> 248  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 248  
gcgtgcgttg tcgtgtcgt t 21

<210> 249  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (2)... (2)  
<223> m5c

<221> modified\_base  
<222> (6)... (6)  
<223> m5c

```

    <221> modified_base
    <222> (10)...(10)
    <223> m5c

    <221> modified_base
    <222> (15)...(15)
    <223> m5c

    <400> 249
cnggcnggc n ggcncgcg                               19

    <210> 250
    <211> 20
    <212> DNA
    <213> Artificial Sequence

    <400> 250
gcggcgggcg gcgcgcgcc                               20

    <210> 251
    <211> 20
    <212> DNA
    <213> Artificial Sequence

    <220>
    <221> modified_base
    <222> (3)...(3)
    <223> I

    <221> modified_base
    <222> (8)...(8)
    <223> I

    <221> modified_base
    <222> (14)...(14)
    <223> I

    <400> 251
agncccgnga acgnattcac                               20

    <210> 252
    <211> 21
    <212> DNA
    <213> Artificial Sequence

    <400> 252
tgtcgtttgt cgttgtcgt t                               21

    <210> 253
    <211> 25
    <212> DNA
    <213> Artificial Sequence

    <400> 253
tgtcgttgtc gttgtcgttg tcgtt                         25

    <210> 254
    <211> 25
    <212> DNA
    <213> Artificial Sequence

```

<400> 254  
tgctcgttgctc gttgctcgttg tcggtt 25  
  
<210> 255  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 255  
tcgtcgtcgtc cggtt 14  
  
<210> 256  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 256  
tgctcgttgctc gtt 13  
  
<210> 257  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 257  
cccccccccc ccccccccccc 20  
  
<210> 258  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 258  
tctagcgttt ttagcgttcc 20  
  
<210> 259  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 259  
tgcaccccc aggccaccat 20  
  
<210> 260  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 260  
tcgtcgtcgtc cgtcgtcgtc gtt 23  
  
<210> 261  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 261  
tcgtcgttggt cgttgctcgtt 20  
  
<210> 262

<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 262  
tcgtcgtttt gtcgttttgt cggt 24

<210> 263  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 263  
tcgtcgttgt cgttttgtcg tt 22

<210> 264  
<211> 39  
<212> DNA  
<213> Artificial Sequence

<400> 264  
ggggagggag gaacttctta aaattccccc agaatgttt 39

<210> 265  
<211> 39  
<212> DNA  
<213> Artificial Sequence

<400> 265  
aaacattctg ggggaatttt aagaagttcc tccctcccc 39

<210> 266  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 266  
atgtttactt cttaaaattc cccagaatg ttt 33

<210> 267  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 267  
aaacattctg ggggaatttt aagaagtaaa cat 33

<210> 268  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 268  
atgtttacta gacaaaattc cccagaatg ttt 33

<210> 269  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 269

39

aaacattctg ggggaatttt gtctagtaaa  
33

cat

<210> 270  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 270  
aaaattgacg ttttaaaaaa

20

<210> 271  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 271  
ccccttgacg ttttcccccc

20

<210> 272  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 272  
ttttcgttgt ttttgtcggt

20

<210> 273  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 273  
tcgtcgtttt gtcgttttgt cggt

24

<210> 274  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 274  
ctgcagcctg ggac

14

<210> 275  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 275  
acccgtcgta attatagtaa aaccc

25

<210> 276  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 276  
ggtacctgtg gggacattgt g

21

<210> 277  
<211> 18

<212> DNA  
<213> Artificial Sequence

<400> 277  
agcaccgaac gtgagagg 18

<210> 278  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 278  
tccatgccgt tcctgccgtt 20

<210> 279  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 279  
tccatgacgg tcctgacggt 20

<210> 280  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 280  
tccatgccgg tcctgccggt 20

<210> 281  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 281  
tccatgcgcg tcctgcgcgt 20

<210> 282  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 282  
ctggtctttc tggttttttt ctgg 24

<210> 283  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 283  
tcaggggtgg ggggaacctt 20

<210> 284  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base

<222> (8)...(8)  
 <223> m5c

<400> 284  
 tccatgangt tcctagttct 20

<210> 285  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<400> 285  
 tccatgatgt tcctagttct 20

<210> 286  
 <211> 26  
 <212> DNA  
 <213> Artificial Sequence

<400> 286  
 cccgaagtca tttcctctta acctgg 26

<210> 287  
 <211> 26  
 <212> DNA  
 <213> Artificial Sequence

<400> 287  
 ccaggttaag aggaaatgac ttcggg 26

<210> 288  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> modified\_base  
 <222> (7)...(7)  
 <223> m5c

<400> 288  
 tcctggnggg gaagt 15

<210> 289  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> modified\_base  
 <222> (2)...(2)  
 <223> m5c

<221> modified\_base  
 <222> (5)...(5)  
 <223> m5c

<221> modified\_base  
 <222> (9)...(9)  
 <223> m5c

<221> modified\_base  
<222> (12)...(12)  
<223> m5c

<221> modified\_base  
<222> (14)...(14)  
<223> m5c

<221> modified\_base  
<222> (16)...(16)  
<223> m5c

<400> 289  
gnggngggng gngngngccc 20

<210> 290  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 290  
tccatgtgct tcctgatgct 20

<210> 291  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 291  
tccatgtcct tcctgatgct 20

<210> 292  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 292  
tccatgtcgt tcctagttct 20

<210> 293  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 293  
tccaagtagt tcctagttct 20

<210> 294  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 294  
tccatgtagt tcctagttct 20

<210> 295  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 295



43

tcccgcgcgt tccgcgcgtt

20

&lt;210&gt; 296

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 296

tcctggcgggt cctggcgggt

20

&lt;210&gt; 297

&lt;211&gt; 15

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 297

tcctggaggg gaagt

15

&lt;210&gt; 298

&lt;211&gt; 15

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 298

tcctgggggg gaagt

15

&lt;210&gt; 299

&lt;211&gt; 15

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 299

tcctggtggg gaagt

15

&lt;210&gt; 300

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 300

tcgtcgtttt gtcgttttgt cggt

24

&lt;210&gt; 301

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 301

ctggtctttc tggttttttt ctgg

24

&lt;210&gt; 302

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 302

tccatgacgt tcctgacgtt

20

&lt;210&gt; 303

&lt;211&gt; 20

&lt;212&gt; DNA

<213> Artificial Sequence

<400> 303  
tccaggactt ctctcaggtt 20

<210> 304  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (2)...(2)  
<223> m5c

<221> modified\_base  
<222> (5)...(5)  
<223> m5c

<221> modified\_base  
<222> (13)...(13)  
<223> m5c

<221> modified\_base  
<222> (21)...(21)  
<223> m5c

<400> 304  
tngtngtttt gtngttttgt ngtt 24

<210> 305  
<211> 29  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 305  
tcgtcgtttt gtcgttttgt cggttttttt 29

<210> 306  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 306  
gctatgacgt tccaaggg 18

<210> 307  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 307  
tcaacgtt 8

<210> 308  
<211> 20

<212> DNA  
<213> Artificial Sequence

<400> 308  
tccaggactt tcctcaggtt 20

<210> 309  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 309  
ctctctgtag gcccgcttgg 20

<210> 310  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 310  
ctttccgttg gaccctggg 20

<210> 311  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 311  
gtccgggcca ggccaaagtc 20

<210> 312  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 312  
gtgcgcgcga gcccgaaatc 20

<210> 313  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (8)...(8)  
<223> I

<221> modified\_base  
<222> (17)...(17)  
<223> I

<400> 313  
tccatgangt tcctgangtt 20

<210> 314  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 314

46

aatagtcgcc ataacaaaac

20

<210> 315  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 315  
aatagtcgcc atggcggggc

20

<210> 316  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_difference  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 316  
tttttccatg tcgttctga tgcttttt

28

<210> 317  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 317  
tcctgtcggtt gaagtttttt

20

<210> 318  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 318  
gctagcttta gagctttaga gctt

24

<210> 319  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 319  
tgctgcttcc cccccccccc

20

<210> 320  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 320  
tcgacgttcc cccccccccc

20

<210> 321  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 321

47

tcgtcgttcc ccccccccc

20

<210> 322  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 322  
tcgtcgttcc ccccccccc

20

<210> 323  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 323  
tcgccgttcc ccccccccc

20

<210> 324  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 324  
tcgtcgatcc ccccccccc

20

<210> 325  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 325  
tcctgacgtt gaagt

15

<210> 326  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 326  
tcctgccgtt gaagt

15

<210> 327  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 327  
tcctgacggt gaagt

15

<210> 328  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 328  
tcctgagctt gaagt

15

<210> 329  
<211> 15  
<212> DNA

<213> Artificial Sequence

<400> 329  
tcctggcggg gaagt 15

<210> 330  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 330  
aaaatctgtg cttttaaaaa a 21

<210> 331  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 331  
gatccagtca cagtacctg gcagaatctg gat 33

<210> 332  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 332  
gatccagatt ctgccaggtc actgtgactg gat 33

<210> 333  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 333  
gatccagtca cagtgactca gcagaatctg gat 33

<210> 334  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<400> 334  
gatccagatt ctgctgagtc actgtgactg gat 33

<210> 335  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (16)...(16)  
<223> m5c

<400> 335  
tcgtcgttcc ccccncccc 20

<210> 336  
<211> 20  
<212> DNA

<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (2)...(2)  
<223> m5c

<221> modified\_base  
<222> (5)...(5)  
<223> m5c

<400> 336  
tngtngttcc cccccccccc 20

<210> 337  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (2)...(2)  
<223> m5c

<400> 337  
tngtcgttcc cccccccccc 20

<210> 338  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (5)...(5)  
<223> m5c

<400> 338  
tcgtngttcc cccccccccc 20

<210> 339  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 339  
tcgtcgtcc cccccccccc 20

<210> 340  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 340  
tcgtcgggtcc cccccccccc 20

<210> 341  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 341  
tcggcggttcc cccccccccc 20

<210> 342  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 342  
ggccttttcc cccccccccc 20

<210> 343  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 343  
tcgtcgtttt gacgttttgt cgtt 24

<210> 344  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 344  
tcgtcgtttt gacgttttga cgtt 24

<210> 345  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 345  
cagtcggttcc cccccccccc 20

<210> 346  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 346  
gcgtcggttcc cccccccccc 20

<210> 347  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 347  
tcgtcattcc cccccccccc 20

<210> 348  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 348  
acgtcggttcc cccccccccc 20

<210> 349  
<211> 20



<212> DNA  
<213> Artificial Sequence

<400> 349  
ctgtcgttcc cccccccccc 20

<210> 350  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 350  
tttttcgtcg ttcccccccc cccc 24

<210> 351  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (18)...(20)  
<223> Biotin moiety attached at 3' end of sequence.

<400> 351  
tcgtcgttcc cccccccccc 20

<210> 352  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (22)...(24)  
<223> Biotin moiety attached at 3' end of sequence.

<400> 352  
tcgtcgtttt gtcgttttgt cggt 24

<210> 353  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 353  
tccagttcct tcctcagtct 20

<210> 354  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (2)...(2)

<223> m5c  
 <400> 354  
 tngtcgtttt gtcgttttgt cggt 24  
 <210> 355  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 355  
 tcctggaggg gaagt 15  
 <210> 356  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 356  
 tcctgaaaag gaagt 15  
 <210> 357  
 <211> 17  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 357  
 tcgtcgttcc cccccc 17  
 <210> 358  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
 <220>  
 <221> modified\_base  
 <222> (2)...(2)  
 <223> m5c  
 <221> modified\_base  
 <222> (5)...(5)  
 <223> m5c  
 <221> modified\_base  
 <222> (13)...(13)  
 <223> m5c  
 <221> modified\_base  
 <222> (21)...(21)  
 <223> m5c  
 <400> 358  
 tngtngtttt gtngttttgt nggt 24  
 <210> 359  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 359  
 ggggtcaagc ttgagggggg 20

<210> 360  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 360  
tgctgcttcc cccccccccc 20  
  
<210> 361  
<211> 14  
<212> DNA  
<213> Artificial Sequence  
  
<400> 361  
tcgtcgtcgt cggt 14  
  
<210> 362  
<211> 14  
<212> DNA  
<213> Artificial Sequence  
  
<400> 362  
tcgtcgtcgt cggt 14  
  
<210> 363  
<211> 14  
<212> DNA  
<213> Artificial Sequence  
  
<400> 363  
tcgtcgtcgt cggt 14  
  
<210> 364  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
  
<400> 364  
tcaacgttga 10  
  
<210> 365  
<211> 8  
<212> DNA  
<213> Artificial Sequence  
  
<400> 365  
tcaacggt 8  
  
<210> 366  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 366  
atagttttcc atttttttac 20  
  
<210> 367  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 367  
aatagtcgcc atcgcgcgac 20  
  
<210> 368  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 368  
aatagtcgcc atcccgggac 20  
  
<210> 369  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 369  
aatagtcgcc atcccccccc 20  
  
<210> 370  
<211> 24  
<212> DNA  
<213> Artificial Sequence  
  
<400> 370  
tgctgctttt gtgcttttgt gctt 24  
  
<210> 371  
<211> 24  
<212> DNA  
<213> Artificial Sequence  
  
<400> 371  
ctgtgctttc tgtgtttttc tgtg 24  
  
<210> 372  
<211> 24  
<212> DNA  
<213> Artificial Sequence  
  
<400> 372  
ctaattcttc taattttttt ctaa 24  
  
<210> 373  
<211> 26  
<212> DNA  
<213> Artificial Sequence  
  
<400> 373  
tcgtcgttgg tgcggttgt gtcgtt 26  
  
<210> 374  
<211> 24  
<212> DNA  
<213> Artificial Sequence  
  
<400> 374  
tcgtcgttgg ttgtcgtttt gggt 24  
  
<210> 375

<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 375  
accatggacg agctgtttcc cctc 24

<210> 376  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 376  
tcgtcgtttt gcgtgcgttt 20

<210> 377  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 377  
ctgtaagtga gcttgagag 20

<210> 378  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 378  
gagaacgctg gaccttcc 18

<210> 379  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 379  
cgggcgactc agtctatcgg 20

<210> 380  
<211> 37  
<212> DNA  
<213> Artificial Sequence

<400> 380  
gttctcagat aaagcggaac cagcaacaga cacagaa 37

<210> 381  
<211> 37  
<212> DNA  
<213> Artificial Sequence

<400> 381  
ttctgtgtct gttgctggtt ccgctttatc tgagaac 37

<210> 382  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 382

56  
23

cagacacaga agcccgatag acg

<210> 383  
<211> 20  
<212> DNA  
<213> Artificial Sequence<400> 383  
agacagacac gaaacgaccg 20<210> 384  
<211> 20  
<212> DNA  
<213> Artificial Sequence<400> 384  
gtctgtccca tgatctcgaa 20<210> 385  
<211> 20  
<212> DNA  
<213> Artificial Sequence<400> 385  
gctggccagc ttacctccg 20<210> 386  
<211> 21  
<212> DNA  
<213> Artificial Sequence<400> 386  
ggggcctcta tacaacctgg g 21<210> 387  
<211> 18  
<212> DNA  
<213> Artificial Sequence<400> 387  
ggggtccctg agactgcc 18<210> 388  
<211> 20  
<212> DNA  
<213> Artificial Sequence<400> 388  
gagaacgctg gaccttccat 20<210> 389  
<211> 20  
<212> DNA  
<213> Artificial Sequence<400> 389  
tccatgtcgg tcctgatgct 20<210> 390  
<211> 20  
<212> DNA

<213> Artificial Sequence  
 <400> 390  
 ctcttgcgac ctggaaggta 20  
 <210> 391  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 391  
 aggtacagcc aggactacga 20  
 <210> 392  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 392  
 accatggacg acctgtttcc cctc 24  
 <210> 393  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 393  
 accatggatt acctttttcc cctt 24  
 <210> 394  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 394  
 atggaaggtc cagcgttctc 20  
 <210> 395  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 395  
 agcatcagga ccgacatgga 20  
 <210> 396  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 396  
 ctctccaagc tcacttacag 20  
 <210> 397  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 397  
 tccctgagac tgccccacct t 21

<210> 398  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 398  
gccaccaaaa cttgtccatg 20

<210> 399  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 399  
gtccatggcg tgcgggatga 20

<210> 400  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 400  
cctctataca acctgggac 19

<210> 401  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 401  
cgggcgactc agtctatcgg 20

<210> 402  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 402  
gcgctaccgg tagcctgagt 20

<210> 403  
<211> 35  
<212> DNA  
<213> Artificial Sequence

<400> 403  
cgactgccga acaggatatc ggtgatcagc actgg 35

<210> 404  
<211> 35  
<212> DNA  
<213> Artificial Sequence

<400> 404  
ccagtgtga tcaccgatat cctgttcggc agtcg 35

<210> 405  
<211> 17  
<212> DNA  
<213> Artificial Sequence



<400> 405  
ccaggttgta tagaggc 17  
  
<210> 406  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 406  
tctcccagcg tacgccat 18  
  
<210> 407  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 407  
tctcccagcg tgcgtttt 18  
  
<210> 408  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 408  
tctcccagcg tgcgccat. 18  
  
<210> 409  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 409  
tctcccgtcg tgcgccat 18  
  
<210> 410  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 410  
ataatcgtag ttcaagcaag 20  
  
<210> 411  
<211> 23  
<212> DNA  
<213> Artificial Sequence  
  
<400> 411  
tcgtcgtttt gtcgttttgt cgt 23  
  
<210> 412  
<211> 24  
<212> DNA  
<213> Artificial Sequence  
  
<400> 412  
tcgtcgtttt gtcgttttgt cggt 24  
  
<210> 413  
<211> 24

<212> DNA  
<213> Artificial Sequence

<400> 413  
tcgtcgtttt gtcgttttgt cggt 24

<210> 414  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_difference  
<222> (3)...(3)  
<223> n is a or c or g or t/u

<221> misc\_difference  
<222> (8)...(8)  
<223> n is a or c or g or t/u

<221> misc\_difference  
<222> (11)...(11)  
<223> n is a or c or g or t/u

<221> misc\_difference  
<222> (16)...(16)  
<223> n is a or c or g or t/u

<221> misc\_difference  
<222> (19)...(19)  
<223> n is a or c or g or t/u

<221> misc\_difference  
<222> (24)...(24)  
<223> n is a or c or g or t/u

<400> 414  
tcttcgtntt ntctgtntnt cgtn 24

<210> 415  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 415  
tctcccagcg tcgcat 17

<210> 416  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 416  
tctcccatcg tcgcat 17

<210> 417  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 417

61

ataatcgtgc gttcaagaaa g

21

<210> 418  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 418  
ataatcgacg ttccccccc

20

<210> 419  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 419  
tctatcgacg ttcaagcaag

20

<210> 420  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 420  
tcctgacggg gagt

14

<210> 421  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 421  
tccatgacgt tcctgatcc

19

<210> 422  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 422  
tccatgacgt tcctgatcc

19

<210> 423  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 423  
tccatgacgt tcctgatcc

19

<210> 424  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 424  
tcctggcgtg gaagt

15

<210> 425  
<211> 19  
<212> DNA

<213> Artificial Sequence  
 <400> 425  
 tccatgacgt tcctgatcc 19  
 <210> 426  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 426  
 tcgtcgctgt tgcgtttct t 21  
 <210> 427  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 427  
 agcagcttta gagctttaga gctt 24  
 <210> 428  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 428  
 cccccccccc ccccccccc cccc 24  
 <210> 429  
 <211> 32  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 429  
 tcgtcgtttt gtcgttttgcg tt 32  
 <210> 430  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 430  
 tcgtcgtttt ttgtcgtttt ttgtcgtt 28  
 <210> 431  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 431  
 tcgtcgtttt tttttttttt 20  
 <210> 432  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
 <400> 432  
 tttttcaacg ttgatttttt 20

<210> 433  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 433  
tttttttttt tttttttttt tttt 24

<210> 434  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 434  
ggggtcgctcg ttttgggggg 20

<210> 435  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 435  
tcgctcgtttt gtcgttttgg gggg 24

<210> 436  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 436  
tcgctcgctgt ctccgcttct tcttgcc 27

<210> 437  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 437  
tcgctcgctgt ctccg 15

<210> 438  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 438  
ctgtaagtga gcttgagag 20

<210> 439  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 439  
gagaacgctg gaccttccat 20

<210> 440  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 440  
ccaggttgta tagaggc 17  
  
<210> 441  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 441  
gctagacggtt agcgtga 17  
  
<210> 442  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 442  
ggagctcttc gaacgccata 20  
  
<210> 443  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 443  
tctccatgat ggttttatcg 20  
  
<210> 444  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 444  
aaggtggggc agtctcaggg a 21  
  
<210> 445  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 445  
atcggaggac tggcgcgccg 20  
  
<210> 446  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 446  
ttaggacaag gtctagggtg 20  
  
<210> 447  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 447  
accacaacga gaggaacgca 20  
  
<210> 448  
<211> 20

<212> DNA  
<213> Artificial Sequence

<400> 448  
ggcagtgcag gctcaccggg 20

<210> 449  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 449  
gaaccttcca tgctgtt 17

<210> 450  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 450  
gctagacgtt agcgtga 17

<210> 451  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 451  
gcttggaggg cctgtaagtg 20

<210> 452  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 452  
gtagccttcc ta 12

<210> 453  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 453  
cggtagcctt ccta 14

<210> 454  
<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 454  
cacggtagcc ttccta 16

<210> 455  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 455  
agcacggtag ccttccta 18

<210> 456  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 456  
gaacgctgga ccttccat 18

<210> 457  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 457  
gaccttccat 10

<210> 458  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 458  
tggaccttcc at 12

<210> 459  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 459  
gctggacctt ccat 14

<210> 460  
<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 460  
acgctggacc ttccat 16

<210> 461  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 461  
taagctctgt caacgccagg 20

<210> 462  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 462  
gagaacgctg gaccttccat gt 22

<210> 463  
<211> 20  
<212> DNA  
<213> Artificial Sequence



<400> 463  
tccatgtcgg tcctgatgct 20  
  
<210> 464  
<211> 21  
<212> DNA  
<213> Artificial Sequence  
  
<400> 464  
ttcatgcctt gcaaaatggc g 21  
  
<210> 465  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 465  
tgctagctgt gcctgtacct 20  
  
<210> 466  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 466  
agcatcagga ccgacatgga 20  
  
<210> 467  
<211> 22  
<212> DNA  
<213> Artificial Sequence  
  
<400> 467  
gaccttccat gtcggtcctg at 22  
  
<210> 468  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 468  
acaaccacga gaacgggaac 20  
  
<210> 469  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 469  
gaaccttcca tgctgttcg 20  
  
<210> 470  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 470  
caatcaatct gaggagaccc 20  
  
<210> 471

<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 471  
tcagctctgg tactttttca 20

<210> 472  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 472  
tggttacggg ctgtcccatg 20

<210> 473  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 473  
gtctatcgga ggactggcgc 20

<210> 474  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 474  
cattttacgg gcgggcgggc 20

<210> 475  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 475  
gaggggacca ttttacgggc 20

<210> 476  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 476  
tgtccagccg aggggaccat 20

<210> 477  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 477  
cgggcttacg gcggatgctg 20

<210> 478  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 478

69

20

tggaccttct atgtcgggtcc

&lt;210&gt; 479

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 479

tgtcccatgt ttttagaagc

20

&lt;210&gt; 480

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 480

gtggttacgg tcgtgcccat

20

&lt;210&gt; 481

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 481

cctccaaatg aaagaccccc

20

&lt;210&gt; 482

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 482

ttgtactctc catgatggtt

20

&lt;210&gt; 483

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 483

ttccatgctg ttccggctgg

20

&lt;210&gt; 484

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 484

gaccttctat gtcggtcctg

20

&lt;210&gt; 485

&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 485

gagaccgctc gaccttcgat

20

&lt;210&gt; 486

&lt;211&gt; 20

&lt;212&gt; DNA

<213> Artificial Sequence

<400> 486  
ttgccccata ttttagaaac 20

<210> 487  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 487  
ttgaaactga ggtgggac 18

<210> 488  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 488  
ctatcggagg actggcgcg c 21

<210> 489  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 489  
cttgaggggc ctcccggcgg 20

<210> 490  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 490  
gctgaacctt ccatgctgtt 20

<210> 491  
<211> 32  
<212> DNA  
<213> Artificial Sequence

<400> 491  
tagaaacagc attcttcttt tagggcagca ca 32

<210> 492  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 492  
agatggttct cagataaagc ggaa 24

<210> 493  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 493  
ttccgcttta tctgagaacc atct 24

<210> 494  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 494  
gtcccagggtt gtatagaggc tgc 23

<210> 495  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 495  
gcgccagtcc tccgatagac 20

<210> 496  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 496  
atcggaggac tggcgcgccg 20

<210> 497  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 497  
ggtctgtccc atatttttag 20

<210> 498  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 498  
tttttcaacg ttgagggggg 20

<210> 499  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 499  
tttttcaagc gttgattttt t 21

<210> 500  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 500  
ggggtcaacg ttgatttttt 20

<210> 501  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 501  
ggggttttca acgttttgag ggggg 25

<210> 502  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 502  
ggttacggtc tgtcccatat 20

<210> 503  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 503  
ctgtcccata tttttagaca 20

<210> 504  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 504  
accatcctga ggccattcgg 20

<210> 505  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 505  
cgtctatcgg gcttctgtgt ctg 23

<210> 506  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 506  
ggccatccca cattgaaagt t 21

<210> 507  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 507  
ccaaatatcg gtggtcaagc ac 22

<210> 508  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 508  
gtgcttgacc accgatattt gg 22

<210> 509  
<211> 26

<212> DNA  
<213> Artificial Sequence

<400> 509  
gtgctgatca ccgatatcct gttcgg 26

<210> 510  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 510  
ggccaaacttt caatgtggga tggcctc 27

<210> 511  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 511  
ttccgccgaa tggcctcagg atggtac 27

<210> 512  
<211> 36  
<212> DNA  
<213> Artificial Sequence

<400> 512  
tatagtccct gagactgccc caccttctca acaacc 36

<210> 513  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 513  
gcagcctcta tacaacctgg gacggga 27

<210> 514  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 514  
ctatcgaggg actggcgcg c 22

<210> 515  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 515  
tatcgaggga ctggcgcgcc g 21

<210> 516  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 516  
gatcgaggga ctggcgcgcc g 21

<210> 517  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<400> 517  
ccgaacagga tatcggatgat cagcac 26

<210> 518  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 518  
ttttgggggc aacgttgagg gggg 24

<210> 519  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 519  
gggggtcaacg ttgagggggg 20

<210> 520  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 520  
cgcgcgcgcg cgcgcgcgcg 20

<210> 521  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 521  
ggggcatgac gttcggggg 20

<210> 522  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 522  
ggggcatgac gttcaaaaaa 20

<210> 523  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 523  
ggggcatgag cttcgggggg 20

<210> 524  
<211> 20  
<212> DNA  
<213> Artificial Sequence



<400> 524  
ggggcatgac gttcggggg 20

<210> 525  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 525  
aaaacatgac gttcaaaaaa 20

<210> 526  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 526  
aaaacatgac gttcggggg 20

<210> 527  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 527  
ggggcatgac gttcaaaaaa 20

<210> 528  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 528  
accatggacg atctgtttcc cctc 24

<210> 529  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 529  
gccatggacg aactgttccc cctc 24

<210> 530  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 530  
cccccccccc ccccccccc 20

<210> 531  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 531  
gggggggggg gggggggggg 20

<210> 532

<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 532  
gctgtaaaaat gaatcggccg 20

<210> 533  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 533  
ttcgggcgga ctctccatt 20

<210> 534  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 534  
tatgccgcgc ccggacttat 20

<210> 535  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 535  
ggggtaatcg atcagggggg 20

<210> 536  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 536  
tttgagaacg ctggaccttc 20

<210> 537  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 537  
gatcgtgat ctaatgctcg 20

<210> 538  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 538  
gtcggtcctg atgctgttcc 20

<210> 539  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 539

tcgtcgtcag ttcgctgtcg 20

<210> 540  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 540  
ctggaccttc catgtcgg 18

<210> 541  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 541  
gctcgttcag cgcgtct 17

<210> 542  
<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 542  
ctggaccttc catgtc 16

<210> 543  
<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 543  
cactgtcctt cgtcga 16

<210> 544  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 544  
cgctggacct tccatgtcgg 20

<210> 545  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 545  
gctgagctca tgccgtctgc 20

<210> 546  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 546  
aacgctggac cttccatgtc 20

<210> 547  
<211> 20  
<212> DNA

<213> Artificial Sequence

<400> 547  
tgcattgccgt acacagctct 20

<210> 548  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 548  
ccttccatgt cggctcctgat 20

<210> 549  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 549  
tactcttcgg atcccttgcg 20

<210> 550  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 550  
ttccatgtcg gtcctgat 18

<210> 551  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 551  
ctgattgctc tctcgtga 18

<210> 552  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 552  
ggcgttattc ctgactcgcc 20

<210> 553  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 553  
cctacgttgt atgcgccag ct 22

<210> 554  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 554  
ggggtaatcg atgagggggg 20

<210> 555  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 555  
ttcgggcgga ctctccatt 20

<210> 556  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 556  
tttttttttt tttttttttt 20

<210> 557  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 557  
gggggttttt tttttggggg 20

<210> 558  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 558  
tttttggggg gggggttttt 20

<210> 559  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 559  
gggggggggg ggggggggt 19

<210> 560  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 560  
aaaaaaaaa aaaaaaaaaa 20

<210> 561  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 561  
cccccaaaaa aaaaaccccc 20

<210> 562  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 562  
aaaaaccccc cccccaaaaa 20

<210> 563  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 563  
tttgaattca ggactggtga ggttgag 27

<210> 564  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 564  
tttgaatcct cagcggctctc cagtggc 27

<210> 565  
<211> 45  
<212> DNA  
<213> Artificial Sequence

<400> 565  
aattctctat cggggcttct gtgtctgttg ctggttccgc tttat 45

<210> 566  
<211> 45  
<212> DNA  
<213> Artificial Sequence

<400> 566  
ctagataaag cggaaccagc aacagacaca gaagccccga tagag 45

<210> 567  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<400> 567  
ttttctagag aggtgcacaa tgctctgg 28

<210> 568  
<211> 29  
<212> DNA  
<213> Artificial Sequence

<400> 568  
tttgaattcc gtgtacagaa gcgagaagc 29

<210> 569  
<211> 31  
<212> DNA  
<213> Artificial Sequence

<400> 569  
tttgcggccg ctagacttaa cctgagagat a 31

<210> 570  
<211> 29

<212> DNA  
<213> Artificial Sequence

<400> 570  
tttgggccca cgagagacag agacacttc 29

<210> 571  
<211> 29  
<212> DNA  
<213> Artificial Sequence

<400> 571  
tttgggcccg cttctcgctt ctgtacacg 29

<210> 572  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 572  
gagaacgctg gaccttccat 20

<210> 573  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 573  
tccatgtcgg tcctgatgct 20

<210> 574  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 574  
ctgtcg 6

<210> 575  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 575  
tcgtga 6

<210> 576  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 576  
cgtcga 6

<210> 577  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 577  
agtgct 6

<210> 578  
<211> 6  
<212> DNA  
<213> Artificial Sequence  
  
<400> 578  
ctgtcg 6  
  
<210> 579  
<211> 6  
<212> DNA  
<213> Artificial Sequence  
  
<400> 579  
agtgct 6  
  
<210> 580  
<211> 6  
<212> DNA  
<213> Artificial Sequence  
  
<400> 580  
cgtcga 6  
  
<210> 581  
<211> 6  
<212> DNA  
<213> Artificial Sequence  
  
<400> 581  
tcgtga 6  
  
<210> 582  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 582  
gagaacgctc cagcttcgat 20  
  
<210> 583  
<211> 17  
<212> DNA  
<213> Artificial Sequence  
  
<400> 583  
gctagacgta agcgtga 17  
  
<210> 584  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 584  
gagaacgctc gaccttccat 20  
  
<210> 585  
<211> 21  
<212> DNA  
<213> Artificial Sequence



<400> 585  
gagaacgctg gacctatcca t 21  
  
<210> 586  
<211> 17  
<212> DNA  
<213> Artificial Sequence  
  
<400> 586  
gctagagggtt agcgtga 17  
  
<210> 587  
<211> 19  
<212> DNA  
<213> Artificial Sequence  
  
<400> 587  
gagaacgctg gacttccat 19  
  
<210> 588  
<211> 17  
<212> DNA  
<213> Artificial Sequence  
  
<400> 588  
tcacgctaac gtctagc 17  
  
<210> 589  
<211> 17  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.  
  
<400> 589  
gctagacgtt agcgtga 17  
  
<210> 590  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 590  
atggaagggtc gagcgttctc 20  
  
<210> 591  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 591  
gagaacgctg gaccttcgat 20  
  
<210> 592  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 592  
gagaacgatg gaccttccat 20  
  
<210> 593  
<211> 17  
<212> DNA  
<213> Artificial Sequence  
  
<400> 593  
gagaacgctg gatccat 17  
  
<210> 594  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 594  
gagaacgctc cagcactgat 20  
  
<210> 595  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 595  
tccatgtcgg tcctgctgat 20  
  
<210> 596  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 596  
atgtcctcgg tcctgatgct 20  
  
<210> 597  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 597  
gagaacgctc caccttccat 20  
  
<210> 598  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 598  
gagaacgctg gaccttcgta 20  
  
<210> 599  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 599  
atggaaggtc cagcgttctc 20

<210> 600  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 600  
tcctga 6

<210> 601  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 601  
tcaacggt 8

<210> 602  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 602  
aacggt 6

<210> 603  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 603  
aacgttga 8

<210> 604  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 604  
tcacgctaac ctctagc 17

<210> 605  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 605  
gagaacgctg gaccttgcac 20

<210> 606  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 606  
gctggacctt ccat 14

<210> 607

<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 607  
gagaacgctg gacatcatcc at 22

<210> 608  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 608  
gagaacgctg gacgctcatc cat 23

<210> 609  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 609  
aacgttgagg ggcac 15

<210> 610  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 610  
atgcccctca acggt 15

<210> 611  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 611  
tcaacgttga 10

<210> 612  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 612  
gctggacctt ccat 14

<210> 613  
<211> 7  
<212> DNA  
<213> Artificial Sequence

<400> 613  
caacgtt 7

<210> 614  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 614

acaacgttga

<210> 615  
<211> 6  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 615

tcacgt

6

<210> 616  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 616

tcaagctt

8

<210> 617  
<211> 6  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 617

tcgtca

6

<210> 618  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 618

aggatatc

8

<210> 619  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 619

tagacgtc

8

<210> 620  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 620

gacgtcat

8

<210> 621  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 621

ccatcgat

8

<210> 622  
<211> 8  
<212> DNA

<213> Artificial Sequence

<400> 622  
atcgatgt 8

<210> 623  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 623  
atgcatgt 8

<210> 624  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 624  
ccatgcat 8

<210> 625  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 625  
agcgctga 8

<210> 626  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 626  
tcagcgct 8

<210> 627  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 627  
ccttcgat 8

<210> 628  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 628  
gtgccggggt ctccgggc 18

<210> 629  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 629  
gctgtggggc ggctcctg 18

<210> 630  
<211> 8  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.  
  
<400> 630  
tcaacggtt 8  
  
<210> 631  
<211> 8  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> FITC moiety attached at 5' end of sequence.  
  
<400> 631  
tcaacggtt 8  
  
<210> 632  
<211> 8  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> FITC moiety attached at 5' end of sequence.  
  
<400> 632  
aacgttga 8  
  
<210> 633  
<211> 7  
<212> DNA  
<213> Artificial Sequence  
  
<400> 633  
tcaacgt 7  
  
<210> 634  
<211> 7  
<212> DNA  
<213> Artificial Sequence  
  
<400> 634  
aacgttg 7  
  
<210> 635  
<211> 6  
<212> DNA  
<213> Artificial Sequence  
  
<400> 635

cgacga

6

<210> 636  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 636

tcaacggt

8

<210> 637  
<211> 5  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 637

tcgga

5

<210> 638  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 638

agaacggt

8

<210> 639  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 639

tcatcgat

8

<210> 640  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 640

taaacggt

8

<210> 641  
<211> 8  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 641

ccaacggt

8

<210> 642  
<211> 6  
<212> DNA  
<213> Artificial Sequence

&lt;400&gt; 642

gctcga

6

<210> 643  
<211> 6  
<212> DNA



<213> Artificial Sequence

<400> 643  
cgacgt 6

<210> 644  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 644  
cgtcgt 6

<210> 645  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 645  
acgtgt 6

<210> 646  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 646  
cgttcg 6

<210> 647  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 647  
gagcaagctg gaccttccat 20

<210> 648  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 648  
cgcgta 6

<210> 649  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 649  
cgtacg 6

<210> 650  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 650  
tcaccggt 8

<210> 651  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 651  
caagagatgc taacaatgca 20

<210> 652  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 652  
acccatcaat agctctgtgc 20

<210> 653  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 653  
ccatcgat 8

<210> 654  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 654  
tcgacgct 8

<210> 655  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 655  
ctagcgct 8

<210> 656  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 656  
taagcgct 8

<210> 657  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 657  
tcgcgaattc gcg 13

<210> 658  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 658  
atggaagggtc cagcgttct 19  
  
<210> 659  
<211> 17  
<212> DNA  
<213> Artificial Sequence  
  
<400> 659  
actggacggtt agcgtga 17  
  
<210> 660  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 660  
cgcctggggc tggctctg 18  
  
<210> 661  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 661  
gtgtcggggt ctccgggc 18  
  
<210> 662  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 662  
gtgccggggt ctccgggc 18  
  
<210> 663  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 663  
cgccgtcgcg gcggttg 18  
  
<210> 664  
<211> 21  
<212> DNA  
<213> Artificial Sequence  
  
<400> 664  
gaagttcacg ttgaggggca t 21  
  
<210> 665  
<211> 21  
<212> DNA  
<213> Artificial Sequence  
  
<400> 665  
atctggtgag ggcaagctat g 21  
  
<210> 666  
<211> 21

<212> DNA  
<213> Artificial Sequence

<400> 666  
gttgaaaccc gagaacatca t 21

<210> 667  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 667  
gcaacggt 8

<210> 668  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 668  
gtaacggt 8

<210> 669  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 669  
cgaacggt 8

<210> 670  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 670  
gaaacggt 8

<210> 671  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 671  
caaacggt 8

<210> 672  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 672  
ctaacggt 8

<210> 673  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 673  
ggaacggt 8

<210> 674  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 674  
tgaacggtt 8

<210> 675  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 675  
acaacggtt 8

<210> 676  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 676  
ttaacggtt 8

<210> 677  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 677  
aaaacggtt 8

<210> 678  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 678  
ataacggtt 8

<210> 679  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 679  
aacgttct 8

<210> 680  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 680  
tccgatcg 8

<210> 681  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 681 tccgtacg	8
<210> 682 <211> 17 <212> DNA <213> Artificial Sequence	
<400> 682 gctagacgct agcgtga	17
<210> 683 <211> 25 <212> DNA <213> Artificial Sequence	
<400> 683 gagaacgctg gacctcatca tccat	25
<210> 684 <211> 20 <212> DNA <213> Artificial Sequence	
<400> 684 gagaacgcta gaccttctat	20
<210> 685 <211> 17 <212> DNA <213> Artificial Sequence	
<400> 685 actagacggt agtgtga	17
<210> 686 <211> 22 <212> DNA <213> Artificial Sequence	
<400> 686 cacaccttg tcaatgtcac gt	22
<210> 687 <211> 22 <212> DNA <213> Artificial Sequence	
<400> 687 tctccatcct atggttttat cg	22
<210> 688 <211> 15 <212> DNA <213> Artificial Sequence	
<400> 688 cgctggacct tccat	15
<210> 689	

<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 689  
caccaccttg gtcaatgtca cgt 23

<210> 690  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 690  
gctagacggt agctgga 17

<210> 691  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 691  
agtgcgattg cagatcg 17

<210> 692  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 692  
ttttcgtttt gtggttttgt gggt 24

<210> 693  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 693  
ttttcgtttg tcgttttgtc gtt 23

<210> 694  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 694  
tttttgtttt gtggttttgt gggt 24

<210> 695  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 695  
accgcatgga ttctaggcca 20

<210> 696  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 696

98

gctagacgtt agcgt

15

<210> 697  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 697  
aacgctggac cttccat

17

<210> 698  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (5)...(5)  
<223> m5c

<400> 698  
tcaangtt

8

<210> 699  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 699  
ccttcgat

8

<210> 700  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 700  
actagacgtt agtgtga

17

<210> 701  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 701  
gctagagggtt agcgtga

17

<210> 702  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 702  
atggactctc cagcgttctc

20

<210> 703  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 703



99

atcgactctc gagcggttctc

20

<210> 704  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 704  
gctagacggt agc

13

<210> 705  
<211> 9  
<212> DNA  
<213> Artificial Sequence

<400> 705  
gctagacgt

9

<210> 706  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 706  
agtgcgattc gagatcg

17

<210> 707  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (5)...(5)  
<223> m5c.

<400> 707  
tcagngct

8

<210> 708  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 708  
ctgattgctc tctcgtga

18

<210> 709  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (2)...(2)  
<223> m5c

<400> 709  
tnaacggt

8

<210> 710

100

<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (6)...(6)  
<223> m5c

<400> 710  
gagaangctg gaccttccat 20

<210> 711  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 711  
gctagacggtt aggctga 17

<210> 712  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 712  
gctacttagc gtga 14

<210> 713  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 713  
gctaccttag cgtga 15

<210> 714  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 714  
atcgacttcg agcgttctc 19

<210> 715  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 715  
atgcactctg cagcgttctc 20

<210> 716  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 716  
agtgactctc cagcgttctc 20

<210> 717

101

<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 717  
gccagatggt agctgga 17

<210> 718  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 718  
atcgactcga gcgttctc 18

<210> 719  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 719  
atcgatcgag cgttctc 17

<210> 720  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)... (3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 720  
gagaacgctc gaccttcgat 20

<210> 721  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 721  
gctagacggt agctgga 17

<210> 722  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 722  
atcgactctc gagcgttctc 20

<210> 723  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 723  
tagacgttag cgtga 15

<210> 724

102

<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 724  
cgactctcga gcgttctc 18

<210> 725  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 725  
ggggtcgacc ttggagggg g 21

<210> 726  
<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 726  
gctaacgtta gcgtga 16

<210> 727  
<211> 9  
<212> DNA  
<213> Artificial Sequence

<400> 727  
cgtcgtcgt 9

<210> 728  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (14)...(14)  
<223> m5c

<400> 728  
gagaacgctg gacnttccat 20

<210> 729  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (18)...(18)  
<223> m5c

<400> 729  
atcgacctac gtgcgtntc 20

<210> 730  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (3)...(3)  
<223> m5c

<400> 730  
atngacctac gtgcgttctc 20

<210> 731  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (7)...(7)  
<223> m5c

<400> 731  
gctagangtt agcgt 15

<210> 732  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (14)...(14)  
<223> m5c

<400> 732  
atcgactctc gagngttctc 20

<210> 733  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 733  
ggggtaatgc atcagggggg 20

<210> 734  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 734  
ggctgtattc ctgactgcc 20

<210> 735  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 735  
ccatgctaac ctctagc 17

<210> 736  
<211> 17

<212> DNA  
<213> Artificial Sequence

<400> 736  
gctagatggt agcgtga 17

<210> 737  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 737  
cgtaccttac ggtga 15

<210> 738  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 738  
tccatgctgg tcctgatgct 20

<210> 739  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 739  
atcgactctc tcgagcgttc tc 22

<210> 740  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 740  
gctagagctt agcgtga 17

<210> 741  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 741  
atcgactctc gagtgttctc 20

<210> 742  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 742  
aacgctcgac cttcgat 17

<210> 743  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 743  
ctcaacgctg gaccttccat 20

<210> 744  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 744  
atcgacctac gtgcgttctc 20

<210> 745  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 745  
gagaatgctg gaccttccat 20

<210> 746  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 746  
tcacgctaac ctctgac 17

<210> 747  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 747  
gagaacgctc cagcactgat 20

<210> 748  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 748  
gagcaagctg gaccttccat 20

<210> 749  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 749  
cgctagaggt tagcgtga 18

<210> 750  
<211> 15

<212> DNA  
<213> Artificial Sequence

<400> 750  
gctagatggt aacgt 15

<210> 751  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 751  
atggaaggtc cacgttctc 19

<210> 752  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 752  
gctagatggt agcgt 15

<210> 753  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 753  
gctagacggt agtgt 15

<210> 754  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 754  
tccatgacgg tcctgatgct 20

<210> 755  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 755  
tccatggcgg tcctgatgct 20

<210> 756  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 756  
gctagacgat agcgt 15

<210> 757  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 757  
gctagtcgat agcgt 15



<210> 758  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 758  
tccatgacgt tcctgatgct 20

<210> 759  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 759  
tccatgtcgt tcctgatgct 20

<210> 760  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (13)...(13)  
<223> m5c

<400> 760  
gctagacggt agngt 15

<210> 761  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 761  
gctaggcggt agcgt 15

<210> 762  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (8)...(8)  
<223> m5c

<400> 762  
tccatgtngg tcctgatgct 20

<210> 763  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (12)...(12)  
<223> m5c

<400> 763  
tccatgtcgg tncatgatgct 20

<210> 764  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (3)...(3)  
<223> m5c

<221> modified\_base  
<222> (10)...(10)  
<223> m5c

<221> modified\_base  
<222> (14)...(14)  
<223> m5c

<400> 764  
atngactctn gagngttctc 20

<210> 765  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 765  
atggaagggtc cagtgttctc 20

<210> 766  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 766  
gcatgacgtt gagct 15

<210> 767  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 767  
gggggtcaacg ttgagggggg 20

<210> 768  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 768  
gggggtcaagt ctgagggggg 20

<210> 769  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 769  
cgcgcgcgcg cgcgcgcgcg 20  
  
<210> 770  
<211> 28  
<212> DNA  
<213> Artificial Sequence  
  
<400> 770  
cccccccccc ccccccccc cccccccc 28  
  
<210> 771  
<211> 35  
<212> DNA  
<213> Artificial Sequence  
  
<400> 771  
cccccccccc ccccccccc ccccccccc ccccc 35  
  
<210> 772  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 772  
tccatgtcgc tcctgacct 20  
  
<210> 773  
<211> 15  
<212> DNA  
<213> Artificial Sequence  
  
<400> 773  
gctaaacgtt agcgt 15  
  
<210> 774  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 774  
tccatgtcga tcctgatgct 20  
  
<210> 775  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 775  
tccatgccgg tcctgatgct 20  
  
<210> 776  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 776  
aaaatcaacg ttgaaaaaaa 20  
  
<210> 777  
<211> 20

<212> DNA  
<213> Artificial Sequence

<400> 777  
tccataacgt tcctgatgct 20

<210> 778  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 778  
tggaggtccc accgagatcg gag 23

<210> 779  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 779  
cgtcgctgctc gtcgctgctc t 21

<210> 780  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 780  
ctgctgctgc tgctgctgct g 21

<210> 781  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 781  
gagaacgctc cgaccttcga t 21

<210> 782  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 782  
gctagatggt agcgt 15

<210> 783  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 783  
gcatgacgtt gagct 15

<210> 784  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature

111

<222> (8)...(10)  
<223> FITC moiety attached at 3' end of sequence.

<400> 784  
tcaatgctga 10

<210> 785  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (8)...(10)  
<223> FITC moiety attached at 3' end of sequence.

<400> 785  
tcaacgttga 10

<210> 786  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (8)...(10)  
<223> Biotin moiety attached at 3' end of sequence.

<400> 786  
tcaacgttga 10

<210> 787  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (8)...(10)  
<223> Biotin moiety attached at 3' end of sequence.

<400> 787  
gcaatattgc 10

<210> 788  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (8)...(10)  
<223> FITC moiety attached at 3' end of sequence.

<400> 788  
gcaatattgc 10

<210> 789  
<211> 10  
<212> DNA

<213> Artificial Sequence

<400> 789  
agttgcaact 10

<210> 790  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 790  
tcttcgaa 8

<210> 791  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 791  
tcaacgtc 8

<210> 792  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 792  
ccatgtcggc cctgatgct 19

<210> 793  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 793  
gtttttatat aatttggg 18

<210> 794  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 794  
tttttgttg tcgttttgc gtt 23

<210> 795  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 795  
ttgggggggg tt 12

<210> 796  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 796  
ggggttggg gtt 13

113

<210> 797  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 797  
ggtggtgtag gttttgg 17

<210> 798  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<221> modified\_base  
<222> (6)...(6)  
<223> m5c

<400> 798  
gagaangctc gaccttcgat 20

<210> 799  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 799  
tcaacgttaa cgtaaagtt 20

<210> 800  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<221> modified\_base  
<222> (8)...(8)  
<223> m5c

<400> 800  
gagcaagntg gaccttccat 20

<210> 801  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<221> modified\_base

114

<222> (6)...(6)  
<223> m5c

<400> 801  
gagaangctc cagcactgat 20

<210> 802  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (5)...(5)  
<223> m5c

<221> misc\_feature  
<222> (8)...(10)  
<223> Biotin moiety attached at 3' end of sequence.

<400> 802  
tcaangttga 10

<210> 803  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (2)...(2)  
<223> m5c

<221> misc\_feature  
<222> (8)...(10)  
<223> Biotin moiety attached at 3' end of sequence.

<400> 803  
gnaatattgc 10

<210> 804  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 804  
tgctgctttt gtcgttttgt gctt 24

<210> 805  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 805  
ctgcgtagc aatttaactg tg 22

<210> 806  
<211> 20  
<212> DNA  
<213> Artificial Sequence



<400> 806  
tccatgacgt tcctgatgct 20  
  
<210> 807  
<211> 28  
<212> DNA  
<213> Artificial Sequence  
  
<400> 807  
tgcattgccgt gcatccgtac acagctct 28  
  
<210> 808  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 808  
tgcattgccgt acacagctct 20  
  
<210> 809  
<211> 12  
<212> DNA  
<213> Artificial Sequence  
  
<400> 809  
tgcattcagct ct 12  
  
<210> 810  
<211> 8  
<212> DNA  
<213> Artificial Sequence  
  
<400> 810  
tgcgctct 8  
  
<210> 811  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 811  
cccccccccc ccccccccc 20  
  
<210> 812  
<211> 12  
<212> DNA  
<213> Artificial Sequence  
  
<400> 812  
cccccccccc cc 12  
  
<210> 813  
<211> 8  
<212> DNA  
<213> Artificial Sequence  
  
<400> 813  
cccccccc 8  
  
<210> 814  
<211> 12

<212> DNA  
<213> Artificial Sequence

<400> 814  
tgcattcagct ct 12

<210> 815  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 815  
tgcatgccgt acacagctct 20

<210> 816  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 816  
gagcaagctg gaccttccat 20

<210> 817  
<211> 32  
<212> DNA  
<213> Artificial Sequence

<400> 817  
tcaacgttaa cgttaacgtt aacgttaacg tt 32

<210> 818  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 818  
gagaacgctc gaccttcgat 20

<210> 819  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 819  
gtccccattt cccagaggag gaaat 25

<210> 820  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 820  
ctagcgctg acgtcatcaa gctag 25

<210> 821  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 821  
ctagcttgat gacgtcagcc gctag 25

<210> 822  
<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 822  
cggctgacgt catcaa 16

<210> 823  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 823  
ctgacgtg 8

<210> 824  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 824  
ctgacgtcat 10

<210> 825  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 825  
attcgatcgg ggcggggcga g 21

<210> 826  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 826  
ctcgccccgc cccgatcgaa t 21

<210> 827  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 827  
gactgacgtc agcgt 15

<210> 828  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<400> 828  
ctagcggctg acgtcataaa gctagc 26

<210> 829  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<400> 829  
ctagctttat gacgtcagcc gctagc 26  
  
<210> 830  
<211> 26  
<212> DNA  
<213> Artificial Sequence  
  
<400> 830  
ctagcggctg agctcataaa gctagc 26  
  
<210> 831  
<211> 25  
<212> DNA  
<213> Artificial Sequence  
  
<400> 831  
ctagtggctg acgtcatcaa gctag 25  
  
<210> 832  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 832  
tccaccacgt ggtctatgct 20  
  
<210> 833  
<211> 24  
<212> DNA  
<213> Artificial Sequence  
  
<400> 833  
gggaatgaaa gattttatta taag 24  
  
<210> 834  
<211> 26  
<212> DNA  
<213> Artificial Sequence  
  
<400> 834  
tctaaaacc atctattctt aaccct 26  
  
<210> 835  
<211> 15  
<212> DNA  
<213> Artificial Sequence  
  
<400> 835  
agctcaacgt catgc 15  
  
<210> 836  
<211> 24  
<212> DNA  
<213> Artificial Sequence  
  
<400> 836  
ttaacggtgg tagcgtatt ggtc 24  
  
<210> 837

<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 837  
ttaagaccaa taccgctacc accg 24

<210> 838  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 838  
gatctagtga tgagtcagcc ggatc 25

<210> 839  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 839  
gatccggctg actcatcact agatc 25

<210> 840  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 840  
tccaagacgt tcctgatgct 20

<210> 841  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 841  
tccatgacgt ccctgatgct 20

<210> 842  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 842  
tccaccacgt ggctgatgct 20

<210> 843  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<400> 843  
ccacgtggac ctctagc 17

<210> 844  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 844

120

tcagaccacg tggtcgggtg ttcctga

27

&lt;210&gt; 845

&lt;211&gt; 27

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 845

tcaggaacac ccgaccacgt ggtctga

27

&lt;210&gt; 846

&lt;211&gt; 18

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 846

catttccacg atttcca

18

&lt;210&gt; 847

&lt;211&gt; 19

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 847

ttcctctctg caagagact

19

&lt;210&gt; 848

&lt;211&gt; 19

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 848

tgtatctctc tgaaggact

19

&lt;210&gt; 849

&lt;211&gt; 25

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 849

ataaagcgaa actagcagca gtttc

25

&lt;210&gt; 850

&lt;211&gt; 25

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 850

gaaactgctg ctagtttcgc tttat

25

&lt;210&gt; 851

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 851

tgcccaaaga ggaaaatttg tttcatacag

30

&lt;210&gt; 852

&lt;211&gt; 30

&lt;212&gt; DNA

121

<213> Artificial Sequence

<400> 852  
ctgtatgaaa caaatTTtcc tctttgggca 30

<210> 853  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 853  
ttagggtag ggtaggggtt 20

<210> 854  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 854  
tccatgagct tcctgatgct 20

<210> 855  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 855  
aaaacatgac gttcaaaaaa 20

<210> 856  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 856  
aaaacatgac gttcgggggg 20

<210> 857  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 857  
ggggcatgag cttcgggggg 20

<210> 858  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 858  
ctaggctgac gtcataagc tagt 24

<210> 859  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<400> 859  
tctgacgtca tctgacgttg gctgacgtct 30

122

<210> 860  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 860  
ggaattagta atagatatag aagtt 25

<210> 861  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<400> 861  
tttacctttt ataaacataa ctaaaacaaa 30

<210> 862  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 862  
gcgttttttt ttgcg 15

<210> 863  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 863  
atatctaadc aaaacattaa caaa 24

<210> 864  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 864  
tctatcccag gtggttcctg ttag 24

<210> 865  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)  
<223> Biotin moiety attached at 5' end of sequence.

<400> 865  
tccatgacgt tctgatgct 20

<210> 866  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (1)...(3)



123

<223> Biotin moiety attached at 5' end of sequence.

<400> 866  
tccatgagct tcctgatgct 20

<210> 867  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (11)...(13)  
<223> FITC moiety attached at 3' end of sequence.

<221> misc\_feature  
<222> (0)...(0)  
<223> Has phosphodiester backbone.

<400> 867  
tttttttttt ttt 13

<210> 868  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_feature  
<222> (11)...(13)  
<223> Biotin moiety attached at 3' end of sequence.

<221> misc\_feature  
<222> (0)...(0)  
<223> Has phosphorothioate and phosphodiester chimeric backbone with phosphodiester on 3' end.

<400> 868  
tttttttttt ttt 13

<210> 869  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 869  
ctagcttgat gagtcagcc gctag 25

<210> 870  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 870  
ttcagttgtc ttgctgctta gctaa 25

<210> 871  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 871 tccatgagct tcctgagtct	20
<210> 872 <211> 25 <212> DNA <213> Artificial Sequence	
<400> 872 ctagcggctg acgtcatcaa tctag	25
<210> 873 <211> 20 <212> DNA <213> Artificial Sequence	
<400> 873 tgctagctgt gcctgtacct	20
<210> 874 <211> 23 <212> DNA <213> Artificial Sequence	
<400> 874 atgctaaagg acgtcacatt gca	23
<210> 875 <211> 23 <212> DNA <213> Artificial Sequence	
<400> 875 tgcaatgtga cgtcctttag cat	23
<210> 876 <211> 31 <212> DNA <213> Artificial Sequence	
<400> 876 gtaggggact ttccgagctc gagatcctat g	31
<210> 877 <211> 31 <212> DNA <213> Artificial Sequence	
<400> 877 cataggatct cgagctcgga aagtccccta c	31
<210> 878 <211> 22 <212> DNA <213> Artificial Sequence	
<400> 878 ctgtcaggaa ctgcaggtaa gg	22
<210> 879 <211> 27	

<212> DNA  
<213> Artificial Sequence

<400> 879  
cataacatag gaatatttac tcctcgc 27

<210> 880  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 880  
ctccagctcc aagaaaggac g 21

<210> 881  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 881  
gaagtctctg gtaagtcttc g 21

<210> 882  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 882  
tgctgctttt gtgcttttgt gctt 24

<210> 883  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 883  
tcgtcgtttt gtggttttgt ggtt 24

<210> 884  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 884  
tcgtcgtttg tcgttttgtc gtt 23

<210> 885  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 885  
tcctgacgtt cggcgcgcgcc cc 22

<210> 886  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 886  
tgctgctttt gtgcttttgt gctt 24

<210> 887  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 887  
tccatgagct tcctgagctt 20

<210> 888  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 888  
tcgtcgtttc gtcgttttga cggt 24

<210> 889  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<400> 889  
tcgtcgtttg cgtgcgtttc gtcgtt 26

<210> 890  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 890  
tcgctgcgt tttgtcgttt tgacgtt 27

<210> 891  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 891  
ttcgtcgttt tgcgttttg tcgtt 25

<210> 892  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 892  
tcctgacggg gaagt 15

<210> 893  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 893  
tcctggcgtg gaagt 15

<210> 894  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 894  
tcctggcggg gaagt 15  
  
<210> 895  
<211> 15  
<212> DNA  
<213> Artificial Sequence  
  
<400> 895  
tcctggcggt gaagt 15  
  
<210> 896  
<211> 15  
<212> DNA  
<213> Artificial Sequence  
  
<400> 896  
tcctgacgtg gaagt 15  
  
<210> 897  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 897  
gcgacgttcg gcgcgcgcc 20  
  
<210> 898  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 898  
gcgacgggcg gcgcgcgcc 20  
  
<210> 899  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 899  
gcggcgtgcg gcgcgcgcc 20  
  
<210> 900  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 900  
gcggcgttcg gcgcgcgcc 20  
  
<210> 901  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 901  
gcgacggtcg gcgcgcgcc 20  
  
<210> 902

<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 902  
gcggcggttcg gcgcgcgccc 20

<210> 903  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 903  
gcgacgtgcg gcgcgcgccc 20

<210> 904  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 904  
tcgtcgtgtg ctccg 15

<210> 905  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 905  
tgtgggggtt ttggttttg 20

<210> 906  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 906  
aggggagggg aggggagggg 20

<210> 907  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 907  
tgtgtgtgtg tgtgtgtgtg t 21

<210> 908  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 908  
ctctctctct ctctctctct ct 22

<210> 909  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 909

129

gggggtcgacg tcgagggggg

20

<210> 910  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 910  
atatatatat atatatatat at

22

<210> 911  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 911  
tttttttttt tttttttttt ttttttt

27

<210> 912  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 912  
tttttttttt tttttttttt t

21

<210> 913  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 913  
tttttttttt ttttttttt

18

<210> 914  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 914  
gctagagggg aggggt

15

<210> 915  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 915  
gctagatggt agggg

15

<210> 916  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 916  
gcatgagggg gagct

15

<210> 917  
<211> 20  
<212> DNA

130

<213> Artificial Sequence

<400> 917  
atggaaggctc cagggggctc 20

<210> 918  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 918  
atggactctg gagggggctc 20

<210> 919  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 919  
atggaaggctc caaggggctc 20

<210> 920  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 920  
gagaaggggg gaccttgat 20

<210> 921  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 921  
gagaaggggg gaccttccat 20

<210> 922  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 922  
gagaaggggc cagcactgat 20

<210> 923  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 923  
tccatgtggg gcctgatgct 20

<210> 924  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 924  
tccatgaggg gcctgatgct 20



131

<210> 925  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 925  
tccatgtggg gcctgctgat 20

<210> 926  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 926  
atggactctc cggggttctc 20

<210> 927  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 927  
atggaaggtc cggggttctc 20

<210> 928  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 928  
atggactctg gaggggtctc 20

<210> 929  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 929  
atggaggctc catggggctc 20

<210> 930  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 930  
atggactctg gggggttctc 20

<210> 931  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 931  
tccatgtggg tggggatgct 20

<210> 932  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 932  
tccatgcggg tgggatgct 20  
  
<210> 933  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 933  
tccatggggg tcctgatgct 20  
  
<210> 934  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 934  
tccatggggt ccctgatgct 20  
  
<210> 935  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 935  
tccatggggt gcctgatgct 20  
  
<210> 936  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 936  
tccatggggt tcctgatgct 20  
  
<210> 937  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 937  
tccatcgggg gcctgatgct 20  
  
<210> 938  
<211> 14  
<212> DNA  
<213> Artificial Sequence  
  
<400> 938  
gctagaggga gtgt 14  
  
<210> 939  
<211> 18  
<212> DNA  
<213> Artificial Sequence  
  
<400> 939  
tttttttttt tttttttt 18  
  
<210> 940  
<211> 21

<212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <221> misc\_difference  
 <222> (2)...(2)  
 <223> m is a or c  
  
 <221> misc\_difference  
 <222> (18)...(18)  
 <223> m is a or c  
  
 <400> 940  
 gmgtcaacg ttgaggmgg g 21  
  
 <210> 941  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence  
  
 <400> 941  
 ggggagttcg ttgagggggg g 21  
  
 <210> 942  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
  
 <400> 942  
 tcgtcgtttc ccccccccc 20  
  
 <210> 943  
 <211> 25  
 <212> DNA  
 <213> Artificial Sequence  
  
 <400> 943  
 ttggggggtt tttttttttt ttttt 25  
  
 <210> 944  
 <211> 23  
 <212> DNA  
 <213> Artificial Sequence  
  
 <400> 944  
 tttaaatttt aaaatttaaa ata 23  
  
 <210> 945  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
  
 <400> 945  
 ttggtttttt tggttttttt ttgg 24  
  
 <210> 946  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
  
 <400> 946

134

tttccctttt ccccttttcc cctc

24

<210> 947  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> misc\_difference  
<222> (21)...(21)  
<223> s is g or c

<400> 947  
ggggtcacgc atgagggggg s

21

<210> 948  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 948  
tccatgacgt tcctgacgtt

20

<210> 949  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 949  
tccatgacgt tcctgacgtt

20

<210> 950  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 950  
tccatgacgt tcctgacgtt

20

<210> 951  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 951  
tccatgacgt tcctgacgtt

20

<210> 952  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 952  
tccatgacgt tcctgacgtt

20

<210> 953  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 953

135

tccatgacgt tcctgacgtt

20

<210> 954  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 954  
tccatgacgt tcctgacgtt

20

<210> 955  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 955  
tccatgacgt tcctgacgtt

20

<210> 956  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 956  
tccatgacgt tcctgacgtt

20

<210> 957  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 957  
tccatgacgt tcctgacgtt

20

<210> 958  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 958  
tccatgacgt tcctgacgtt

20

<210> 959  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 959  
gggggacgat cgtcggggg

19

<210> 960  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 960  
gggggtcgta cgacggggg

20

<210> 961  
<211> 24  
<212> DNA

<213> Artificial Sequence

<400> 961  
tttttttttt tttttttttt tttt 24

<210> 962  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 962  
aaaaaaaaaa aaaaaaaaaa aaaa 24

<210> 963  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 963  
cccccccccc ccccccccc cccc 24

<210> 964  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 964  
tcgtcgtttt gtcgttttgt cgtt 24

<210> 965  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 965  
tcgtcgtttt gtcgttttgt cgtt 24

<210> 966  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 966  
tcgtcgtttt gtcgttttgt cgtt 24

<210> 967  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 967  
tcgtcgtttt gtcgttttgt cgtt 24

<210> 968  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 968  
ggggtcaacg ttgagggggg 20

<210> 969  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 969  
gggggtcaacg ttgagggggg 20

<210> 970  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 970  
gggggtcaagc ttgagggggg 20

<210> 971  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 971  
tgctgcttcc ccccccccc 20

<210> 972  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 972  
ggggacgtcg acgtgggggg 20

<210> 973  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 973  
gggggtcgtcg acgagggggg 20

<210> 974  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 974  
gggggtcgacg tacgtcgagg gggg 24

<210> 975  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 975  
ggggaccggt accggtgggg gg 22

<210> 976  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 976  
gggtcgacgt cgagggggg 19  
  
<210> 977  
<211> 19  
<212> DNA  
<213> Artificial Sequence  
  
<400> 977  
ggggtcgacg tcgaggggg 19  
  
<210> 978  
<211> 22  
<212> DNA  
<213> Artificial Sequence  
  
<400> 978  
ggggaacgtt aacgttggg gg 22  
  
<210> 979  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 979  
ggggtcaccg gtgagggggg 20  
  
<210> 980  
<211> 22  
<212> DNA  
<213> Artificial Sequence  
  
<400> 980  
ggggtcgttc gaacgagggg gg 22  
  
<210> 981  
<211> 22  
<212> DNA  
<213> Artificial Sequence  
  
<400> 981  
ggggacgttc gaacgtggg gg 22  
  
<210> 982  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
  
<400> 982  
tcaacttga 10  
  
<210> 983  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
  
<400> 983  
tcaagcttga 10  
  
<210> 984  
<211> 12



<212> DNA  
<213> Artificial Sequence

<400> 984  
tcacgatcgt ga 12

<210> 985  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 985  
tcagcatgct ga 12

<210> 986  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 986  
gggggagcat gctggggggg 20

<210> 987  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 987  
gggggggggg gggggggggg 20

<210> 988  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 988  
gggggacgat atcgtcggg gg 22

<210> 989  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 989  
gggggacgac gtcgtcggg gg 22

<210> 990  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 990  
gggggacgag ctcgtcggg gg 22

<210> 991  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 991  
gggggacgta cgtcggggg 20

<210> 992  
<211> 8  
<212> DNA  
<213> Artificial Sequence  
  
<400> 992  
tcaacggt 8  
  
<210> 993  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 993  
tccataccgg tcctgatgct 20  
  
<210> 994  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 994  
tccataccgg tcctaccggt 20  
  
<210> 995  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 995  
gggggacgat cgttgggggg 20  
  
<210> 996  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 996  
ggggaacgat cgtcgggggg 20  
  
<210> 997  
<211> 21  
<212> DNA  
<213> Artificial Sequence  
  
<400> 997  
ggggggacga tcgtcggggg g 21  
  
<210> 998  
<211> 21  
<212> DNA  
<213> Artificial Sequence  
  
<400> 998  
gggggacgat cgtcgggggg g 21  
  
<210> 999  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 999  
aaagacgtta aa 12

<210> 1000  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1000  
aaagagctta aa 12

<210> 1001  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (6)...(6)  
<223> m5c

<400> 1001  
aaagangtta aa 12

<210> 1002  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1002  
aaattcggaa aa 12

<210> 1003  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1003  
gggggtcatc gatgagggg g 21

<210> 1004  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1004  
gggggtcaac gttgagggg g 21

<210> 1005  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1005  
atgtagctta ataacaaagc 20

<210> 1006  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1006  
ggatcccttg agttacttct 20

<210> 1007  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1007  
ccattccact tctgattacc 20

<210> 1008  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1008  
tatgtattat catgtagata 20

<210> 1009  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1009  
agcctacgta ttcaccctcc 20

<210> 1010  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1010  
ttcctgcaac tactattgta 20

<210> 1011  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1011  
atagaaggcc ctacaccagt 20

<210> 1012  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1012  
ttacaccggt ctatggaggt 20

<210> 1013  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1013  
ctaaccagat caagtctagg 20

<210> 1014

<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1014  
cctagacttg atctggtag 20

<210> 1015  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1015  
tataagcctc gtccgacatg 20

<210> 1016  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1016  
catgtcggac gaggcttata 20

<210> 1017  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1017  
tggtggtggg gagtaagctc 20

<210> 1018  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1018  
gagctactcc cccaccacca 20

<210> 1019  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1019  
gccttcgatc ttcggtggga 20

<210> 1020  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1020  
tggaattctc ttgcccgtct 20

<210> 1021  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1021

144

atgctgtagc ccagcgataa 20

<210> 1022  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1022  
accgaatcag cggaaagtga 20

<210> 1023  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1023  
tccatgacgt tcctgacgtt 20

<210> 1024  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 1024  
ggagaaacc atgagctcat ctgg 24

<210> 1025  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1025  
accacagacc agcaggcaga 20

<210> 1026  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1026  
gagcgtgaac tgcgcgaaga 20

<210> 1027  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1027  
tcggtaccct tgcagcggtt 20

<210> 1028  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1028  
ctggagccct agccaaggat 20

<210> 1029  
<211> 20  
<212> DNA

<213> Artificial Sequence

<400> 1029  
gcgactccat caccagcgat 20

<210> 1030  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1030  
cctgaagtaa gaaccagatg t 21

<210> 1031  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1031  
ctgtgttatc tgacatacac c 21

<210> 1032  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1032  
aattagcctt aggtgattgg g 21

<210> 1033  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1033  
acatctgggtt cttacttcag g 21

<210> 1034  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 1034  
ataagtcata ttttggaac tac 23

<210> 1035  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1035  
cccaatcacc taaggctaat t 21

<210> 1036  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1036  
ggggtcgtcg acgagggggg 20

146

<210> 1037  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 1037  
ggggtcgttc gaacgagggg gg 22

<210> 1038  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 1038  
ggggacgttc gaacgtgggg gg 22

<210> 1039  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (9)...(9)  
<223> n is 5-methylcytosine.

<400> 1039  
tcctggcgng gaagt 15

<210> 1040  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 1040  
ggggaacgac gtcgttgggg gg 22

<210> 1041  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1041  
ggggaacgta cgtcgggggg 20

<210> 1042  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 1042  
ggggaacgta cgtacgttg gggg 24

<210> 1043  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1043  
ggggtcaccg gtgagggggg 20



147

<210> 1044  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 1044  
gggggtcgacg tacgtcgagg gggg 24

<210> 1045  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 1045  
ggggaccggt accggtgggg gg 22

<210> 1046  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 1046  
gggtcgacgt cgagggggg 19

<210> 1047  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 1047  
ggggtcgacg tcgagggg 18

<210> 1048  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<400> 1048  
ggggaacgtt aacgttgggg gg 22

<210> 1049  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 1049  
ggggacgtcg acgtggggg 19

<210> 1050  
<211> 34  
<212> DNA  
<213> Artificial Sequence

<400> 1050  
gcactcttcg aagctacagc cggcagcctc tgat 34

<210> 1051  
<211> 32  
<212> DNA  
<213> Artificial Sequence

148

<400> 1051  
cggctcttcc atgaggtctt tgctaattctt gg 32

<210> 1052  
<211> 35  
<212> DNA  
<213> Artificial Sequence

<400> 1052  
cggctcttcc atgaaagtct ttggacgatg tgagc 35

<210> 1053  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1053  
tcctgcaggt taagt 15

<210> 1054  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1054  
gggggtcgtt cgttgggggg 20

<210> 1055  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1055  
gggggatgat tggtgggggg 20

<210> 1056  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> modified\_base  
<222> (7)...(7)  
<223> m5c

<221> modified\_base  
<222> (11)...(11)  
<223> m5c

<400> 1056  
gggggangat nggtgggggg 20

<210> 1057  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1057  
gggggagcta gcttgggggg 20

<210> 1058

149

<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1058  
ggttcttttg gtccttgtct 20

<210> 1059  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1059  
ggttcttttg gtcctcgtct 20

<210> 1060  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1060  
ggttcttttg gtccttatct 20

<210> 1061  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1061  
ggttcttggg ttccttgtct 20

<210> 1062  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1062  
tggctctttg gtccttgtct 20

<210> 1063  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1063  
ggttcaaag gtccttgtct 20

<210> 1064  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1064  
gggtcttttg ggccttgtct 20

<210> 1065  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 1065

150

tccaggactt ctctcaggtt tttt 24

<210> 1066  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1066  
tccaaaactt ctctcaaatt 20

<210> 1067  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 1067  
tactactttt atacttttat actt 24

<210> 1068  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<400> 1068  
tgtgtgtgtg tgtgtgtgtg tgtg 24

<210> 1069  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<400> 1069  
ttgttggtgt tggttggtgt tggtg 25

<210> 1070  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 1070  
ggctccgggg aggggaatttt tgtctat 27

<210> 1071  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<400> 1071  
gggacgatcg tcggggggg 19

<210> 1072  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1072  
gggtcgtcga cgaggggggg 20

<210> 1073  
<211> 19  
<212> DNA

151

<213> Artificial Sequence

<400> 1073  
ggtcgctcgac gagggggggg 19

<210> 1074  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1074  
gggtcgtcgt cgtggggggg 20

<210> 1075  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1075  
ggggacgatc gtcggggggg 20

<210> 1076  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<400> 1076  
ggggacgtcg tcgtgggggg 20

<210> 1077  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<400> 1077  
ggggtcgacg tcgacgtcga gggggggg 27

<210> 1078  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1078  
ggggaaccgc ggttgggggg g 21

<210> 1079  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<400> 1079  
ggggacgacg tcgtgggggg g 21

<210> 1080  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<400> 1080  
tcgtcgtcgt cgtcgtgggg ggg 23

152

<210> 1081  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1081  
tcctgccggg gaagt 15

<210> 1082  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1082  
tcctgcaggg gaagt 15

<210> 1083  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1083  
tcctgaaggg gaagt 15

<210> 1084  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1084  
tcctggcggg caagt 15

<210> 1085  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1085  
tcctggcggg taagt 15

<210> 1086  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1086  
tcctggcggg aaagt 15

<210> 1087  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1087  
tccgggcggg gaagt 15

<210> 1088  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<400> 1088  
tcggggcggg gaagt 15  
  
<210> 1089  
<211> 15  
<212> DNA  
<213> Artificial Sequence  
  
<400> 1089  
tcccggcggg gaagt 15  
  
<210> 1090  
<211> 15  
<212> DNA  
<213> Artificial Sequence  
  
<400> 1090  
gggggacgtt ggggg 15  
  
<210> 1091  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 1091  
ggggtttttt ttttggggg 20  
  
<210> 1092  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
  
<400> 1092  
ggggccccc ccccggggg 20  
  
<210> 1093  
<211> 21  
<212> DNA  
<213> Artificial Sequence  
  
<400> 1093  
ggggttgttg ttgtggggg g 21  
  
<210> 1094  
<211> 30  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<223> Synthetic Sequence  
  
<400> 1094  
tttttttttt tttttttttt tttttttttt 30  
  
<210> 1095  
<211> 30  
<212> DNA  
<213> Artificial Sequence  
  
<220>  
<223> Synthetic Sequence

<400> 1095  
aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 30

<210> 1096  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Sequence

<400> 1096  
cccccccccc ccccccccccc ccccccccccc 30

<210> 1097  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Sequence

<400> 1097  
cgcgcgcgcg cgcgcgcgcg cgcgcgcgcg 30

<210> 1098  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Sequence

<400> 1098  
gattttatcg tc 12

<210> 1099  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1099  
tcgatttttc ga 12

<210> 1100  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1100  
tcatttttat ga 12

<210> 1101  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1101  
gttttttacg ac 12



155

<210> 1102  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1102  
tcaatttttt ga 12

<210> 1103  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1103  
acgttttttac gt 12

<210> 1104  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1104  
tcgttttttac ga 12

<210> 1105  
<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 1105  
tcgattttta cgtcga 16

<210> 1106  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 1106  
aatttttttaa cggt 14

<210> 1107  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 1107  
tcgtttttta acga 14

<210> 1108  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 1108  
acgtttttta acgt 14

<210> 1109  
<211> 13  
<212> DNA  
<213> Artificial Sequence

<400> 1109 gatttttatc gtc	13
<210> 1110 <211> 14 <212> DNA <213> Artificial Sequence	
<400> 1110 gacgattttt cgtc	14
<210> 1111 <211> 14 <212> DNA <213> Artificial Sequence	
<400> 1111 gatttttagct cgtc	14
<210> 1112 <211> 12 <212> DNA <213> Artificial Sequence	
<400> 1112 gatttttacg tc	12
<210> 1113 <211> 10 <212> DNA <213> Artificial Sequence	
<400> 1113 attttatcgt	10
<210> 1114 <211> 14 <212> DNA <213> Artificial Sequence	
<400> 1114 aacgattttt cgtt	14
<210> 1115 <211> 12 <212> DNA <213> Artificial Sequence	
<400> 1115 tcacttttgt ga	12
<210> 1116 <211> 10 <212> DNA <213> Artificial Sequence	
<400> 1116 tcgtatttta	10
<210> 1117 <211> 14	

<212> DNA  
<213> Artificial Sequence

<400> 1117  
acttttgtac cggt 14

<210> 1118  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 1118  
tcgatttttc gacgtcga 18

<210> 1119  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1119  
acgatttttc gt 12

<210> 1120  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1120  
gatgatcgtc 10

<210> 1121  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1121  
tcgatgtcga 10

<210> 1122  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1122  
tcatgtatga 10

<210> 1123  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1123  
gtgttacgac 10

<210> 1124  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1124  
tcaatgttga 10

<210> 1125  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1125  
acgtgtacgt 10

<210> 1126  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1126  
tcgtgtacga 10

<210> 1127  
<211> 14  
<212> DNA  
<213> Artificial Sequence

<400> 1127  
tcgatgtacg tcga 14

<210> 1128  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1128  
aatgttaacg tt 12

<210> 1129  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1129  
tcgtgttaac ga 12

<210> 1130  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1130  
acgtgttaac gt 12

<210> 1131  
<211> 11  
<212> DNA  
<213> Artificial Sequence

<400> 1131  
gatgtatcgt c 11

<210> 1132  
<211> 12  
<212> DNA  
<213> Artificial Sequence

<400> 1132 gacgatgtcg tc	12
<210> 1133 <211> 12 <212> DNA <213> Artificial Sequence	
<400> 1133 gatgagctcg tc	12
<210> 1134 <211> 10 <212> DNA <213> Artificial Sequence	
<400> 1134 gatgtacgtc	10
<210> 1135 <211> 8 <212> DNA <213> Artificial Sequence	
<400> 1135 atgatcgt	8
<210> 1136 <211> 12 <212> DNA <213> Artificial Sequence	
<400> 1136 aacgatgtcg tt	12
<210> 1137 <211> 10 <212> DNA <213> Artificial Sequence	
<400> 1137 tcactggtga	10
<210> 1138 <211> 8 <212> DNA <213> Artificial Sequence	
<400> 1138 tcgtatga	8
<210> 1139 <211> 12 <212> DNA <213> Artificial Sequence	
<400> 1139 actggtaccg gt	12
<210> 1140	

160

<211> 16  
<212> DNA  
<213> Artificial Sequence

<400> 1140  
tcgatgtcga cgtcga 16

<210> 1141  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<400> 1141  
acgatgtcgt 10

<210> 1142  
<211> 31  
<212> DNA  
<213> Artificial Sequence

<400> 1142  
tgcaggaagt ccgggttttc cccaaccccc c 31

<210> 1143  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic Sequence

<400> 1143  
gacggt 6

<210> 1144  
<211> 6  
<212> DNA  
<213> Artificial Sequence

<400> 1144  
gtcgtt 6

<210> 1145  
<211> 8  
<212> DNA  
<213> Artificial Sequence

<400> 1145  
tcgtcgtt 8